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(54) Title: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF LUNG CANCER AND METHODS FOR THEIR USE

(57) Abstract

Compounds and methods for treating lung cancer are provided. The inventive compounds include polypeptides containing at least a portion of a lung tumor protein. Vaccines and pharmaceutical compositions for immunotherapy of lung cancer comprising such polypeptides, or polynucleotides encoding such polypeptides, are also provided, together with polynucleotides for preparing the inventive polypeptides.

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COMPOUNDS FOR THERAPY AND DIAGNOSIS
OF LUNG CANCER AND METHODS FOR THEIR USE

5   TECHNICAL FIELD

The present invention relates generally to compositions and methods for the treatment of lung cancer. The invention is more specifically related to nucleotide sequences that are preferentially expressed in lung tumor tissue, together with polypeptides encoded by such nucleotide sequences. The inventive nucleotide sequences and polypeptides may be used 10   in vaccines and pharmaceutical compositions for the treatment of lung cancer.

BACKGROUND OF THE INVENTION

Lung cancer is the primary cause of cancer death among both men and women in the U.S., with an estimated 172,000 new cases being reported in 1994. The five-year 15   survival rate among all lung cancer patients, regardless of the stage of disease at diagnosis, is only 13%. This contrasts with a five-year survival rate of 46% among cases detected while the disease is still localized. However, only 16% of lung cancers are discovered before the disease has spread.

Early detection is difficult since clinical symptoms are often not seen until the 20   disease has reached an advanced stage. Currently, diagnosis is aided by the use of chest x-rays, analysis of the type of cells contained in sputum and fiberoptic examination of the bronchial passages. Treatment regimens are determined by the type and stage of the cancer, and include surgery, radiation therapy and/or chemotherapy. In spite of considerable research into therapies for the disease, lung cancer remains difficult to treat.

25   Accordingly, there remains a need in the art for improved vaccines, treatment methods and diagnostic techniques for lung cancer.

SUMMARY OF THE INVENTION

Briefly stated, the present invention provides compounds and methods for the 30   therapy of lung cancer. In a first aspect, isolated polynucleotides encoding lung tumor polypeptides are provided, such polynucleotides comprising a nucleotide sequence selected

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herein; and (b) detecting in the sample a protein or polypeptide that binds to the binding agent. In preferred embodiments, the binding agent is an antibody, most preferably a monoclonal antibody.

In related aspects, methods are provided for monitoring the progression of lung cancer in a patient, comprising: (a) contacting a biological sample obtained from a patient with a binding agent that is capable of binding to one of the polypeptides disclosed herein; (b) determining in the sample an amount of a protein or polypeptide that binds to the binding agent; (c) repeating steps (a) and (b); and comparing the amounts of polypeptide detected in steps (b) and (c).

Within related aspects, the present invention provides antibodies, preferably monoclonal antibodies, that bind to the inventive polypeptides, as well as diagnostic kits comprising such antibodies, and methods of using such antibodies to inhibit the development of lung cancer.

The present invention further provides methods for detecting lung cancer comprising: (a) obtaining a biological sample from a patient; (b) contacting the sample with a first and a second oligonucleotide primer in a polymerase chain reaction, at least one of the oligonucleotide primers being specific for a polynucleotide that encodes one of the polypeptides disclosed herein; and (c) detecting in the sample a DNA sequence that amplifies in the presence of the first and second oligonucleotide primers. In a preferred embodiment, at least one of the oligonucleotide primers comprises at least about 10 contiguous nucleotides of a polynucleotide comprising a sequence selected from the group consisting of SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181.

In a further aspect, the present invention provides a method for detecting lung cancer in a patient comprising: (a) obtaining a biological sample from the patient; (b) contacting the sample with an oligonucleotide probe specific for a polynucleotide that encodes one of the polypeptides disclosed herein; and (c) detecting in the sample a DNA sequence that hybridizes to the oligonucleotide probe. Preferably, the oligonucleotide probe comprises at least about 15 contiguous nucleotides of a polynucleotide comprising a sequence selected from the group consisting of SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181.In related aspects, diagnostic kits comprising the above oligonucleotide probes or primers are provided.

5

SEQ ID NO: 14 is the determined cDNA sequence for L355C1.cons

SEQ ID NO: 15 is the determined cDNA sequence for L366C1.cons

SEQ ID NO: 16 is the determined cDNA sequence for L163C1a

SEQ ID NO: 17 is the determined cDNA sequence for LT86-1

5   SEQ ID NO: 18 is the determined cDNA sequence for LT86-2

SEQ ID NO: 19 is the determined cDNA sequence for LT86-3

SEQ ID NO: 20 is the determined cDNA sequence for LT86-4

SEQ ID NO: 21 is the determined cDNA sequence for LT86-5

SEQ ID NO: 22 is the determined cDNA sequence for LT86-6

10   SEQ ID NO: 23 is the determined cDNA sequence for LT86-7

SEQ ID NO: 24 is the determined cDNA sequence for LT86-8

SEQ ID NO: 25 is the determined cDNA sequence for LT86-9

SEQ ID NO: 26 is the determined cDNA sequence for LT86-10

SEQ ID NO: 27 is the determined cDNA sequence for LT86-11

15   SEQ ID NO: 28 is the determined cDNA sequence for LT86-12

SEQ ID NO: 29 is the determined cDNA sequence for LT86-13

SEQ ID NO: 30 is the determined cDNA sequence for LT86-14

SEQ ID NO: 31 is the determined cDNA sequence for LT86-15

SEQ ID NO: 32 is the predicted amino acid sequence for LT86-1

20   SEQ ID NO: 33 is the predicted amino acid sequence for LT86-2

SEQ ID NO: 34 is the predicted amino acid sequence for LT86-3

SEQ ID NO: 35 is the predicted amino acid sequence for LT86-4

SEQ ID NO: 36 is the predicted amino acid sequence for LT86-5

SEQ ID NO: 37 is the predicted amino acid sequence for LT86-6

25   SEQ ID NO: 38 is the predicted amino acid sequence for LT86-7

SEQ ID NO: 39 is the predicted amino acid sequence for LT86-8

SEQ ID NO: 40 is the predicted amino acid sequence for LT86-9

SEQ ID NO: 41 is the predicted amino acid sequence for LT86-10

SEQ ID NO: 42 is the predicted amino acid sequence for LT86-11

30   SEQ ID NO: 43 is the predicted amino acid sequence for LT86-12

SEQ ID NO: 74 is the predicted amino acid sequence for LT86-21

SEQ ID NO: 75 is the predicted amino acid sequence for LT86-22

SEQ ID NO: 76 is the predicted amino acid sequence for LT86-26

SEQ ID NO: 77 is the predicted amino acid sequence for LT86-27

5   SEQ ID NO: 78 is the determined extended cDNA sequence for L86S-12

SEQ ID NO: 79 is the determined extended cDNA sequence for L86S-36

SEQ ID NO: 80 is the determined extended cDNA sequence for L86S-46

SEQ ID NO: 81 is the predicted extended amino acid sequence for L86S-12

SEQ ID NO: 82 is the predicted extended amino acid sequence for L86S-36

10   SEQ ID NO: 83 is the predicted extended amino acid sequence for L86S-46

SEQ ID NO: 84 is the determined 5'cDNA sequence for L86S-6

SEQ ID NO: 85 is the determined 5'cDNA sequence for L86S-11

SEQ ID NO: 86 is the determined 5'cDNA sequence for L86S-14

SEQ ID NO: 87 is the determined 5'cDNA sequence for L86S-29

15   SEQ ID NO: 88 is the determined 5'cDNA sequence for L86S-34

SEQ ID NO: 89 is the determined 5'cDNA sequence for L86S-39

SEQ ID NO: 90 is the determined 5'cDNA sequence for L86S-47

SEQ ID NO: 91 is the determined 5'cDNA sequence for L86S-49

SEQ ID NO: 92 is the determined 5'cDNA sequence for L86S-51

20   SEQ ID NO: 93 is the predicted amino acid sequence for L86S-6

SEQ ID NO: 94 is the predicted amino acid sequence for L86S-11

SEQ ID NO: 95 is the predicted amino acid sequence for L86S-14

SEQ ID NO: 96 is the predicted amino acid sequence for L86S-29

SEQ ID NO: 97 is the predicted amino acid sequence for L86S-34

25   SEQ ID NO: 98 is the predicted amino acid sequence for L86S-39

SEQ ID NO: 99 is the predicted amino acid sequence for L86S-47

SEQ ID NO: 100 is the predicted amino acid sequence for L86S-49

SEQ ID NO: 101 is the predicted amino acid sequence for L86S-51

SEQ ID NO: 102 is the determined DNA sequence for SLT-T1

30   SEQ ID NO: 103 is the determined 5' cDNA sequence for SLT-T2

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SEQ ID NO: 134 is the determined cDNA sequence for PSLT-69

SEQ ID NO: 135 is the determined cDNA sequence for PSLT-71

SEQ ID NO: 136 is the determined cDNA sequence for PSLT-73

SEQ ID NO: 137 is the determined cDNA sequence for PSLT-79

5   SEQ ID NO: 138 is the determined cDNA sequence for PSLT-03

SEQ ID NO: 139 is the determined cDNA sequence for PSLT-09

SEQ ID NO: 140 is the determined cDNA sequence for PSLT-011

SEQ ID NO: 141 is the determined cDNA sequence for PSLT-041

SEQ ID NO: 142 is the determined cDNA sequence for PSLT-62

10   SEQ ID NO: 143 is the determined cDNA sequence for PSLT-6

SEQ ID NO: 144 is the determined cDNA sequence for PSLT-37

SEQ ID NO: 145 is the determined cDNA sequence for PSLT-74

SEQ ID NO: 146 is the determined cDNA sequence for PSLT-010

SEQ ID NO: 147 is the determined cDNA sequence for PSLT-012

15   SEQ ID NO: 148 is the determined cDNA sequence for PSLT-037

SEQ ID NO: 149 is the determined 5' cDNA sequence for SAL-3

SEQ ID NO: 150 is the determined 5' cDNA sequence for SAL-24

SEQ ID NO: 151 is the determined 5' cDNA sequence for SAL-25

SEQ ID NO: 152 is the determined 5' cDNA sequence for SAL-33

20   SEQ ID NO: 153 is the determined 5' cDNA sequence for SAL-50

SEQ ID NO: 154 is the determined 5' cDNA sequence for SAL-57

SEQ ID NO: 155 is the determined 5' cDNA sequence for SAL-66

SEQ ID NO: 156 is the determined 5' cDNA sequence for SAL-82

SEQ ID NO: 157 is the determined 5' cDNA sequence for SAL-99

25   SEQ ID NO: 158 is the determined 5' cDNA sequence for SAL-104

SEQ ID NO: 159 is the determined 5' cDNA sequence for SAL-109

SEQ ID NO: 160 is the determined 5' cDNA sequence for SAL-5

SEQ ID NO: 161 is the determined 5' cDNA sequence for SAL-8

SEQ ID NO: 162 is the determined 5' cDNA sequence for SAL-12

30   SEQ ID NO: 163 is the determined 5' cDNA sequence for SAL-14

SEQ ID NO: 194 is the predicted amino acid sequence for SAL-5

SEQ ID NO: 195 is the predicted amino acid sequence for SAL-8

SEQ ID NO: 196 is the predicted amino acid sequence for SAL-12

SEQ ID NO: 197 is the predicted amino acid sequence for SAL-14

5   SEQ ID NO: 198 is the predicted amino acid sequence for SAL-16

SEQ ID NO: 199 is the predicted amino acid sequence for SAL-23

SEQ ID NO: 200 is the predicted amino acid sequence for SAL-26

SEQ ID NO: 201 is the predicted amino acid sequence for SAL-29

SEQ ID NO: 202 is the predicted amino acid sequence for SAL-32

10   SEQ ID NO: 203 is the predicted amino acid sequence for SAL-39

SEQ ID NO: 204 is the predicted amino acid sequence for SAL-42

SEQ ID NO: 205 is the predicted amino acid sequence for SAL-43

SEQ ID NO: 206 is the predicted amino acid sequence for SAL-44

SEQ ID NO: 207 is the predicted amino acid sequence for SAL-48

15   SEQ ID NO: 208 is the predicted amino acid sequence for SAL-68

SEQ ID NO: 209 is the predicted amino acid sequence for SAL-72

SEQ ID NO: 210 is the predicted amino acid sequence for SAL-77

SEQ ID NO: 211 is the predicted amino acid sequence for SAL-86

SEQ ID NO: 212 is the predicted amino acid sequence for SAL-88

20   SEQ ID NO: 213 is the predicted amino acid sequence for SAL-93

SEQ ID NO: 214 is the predicted amino acid sequence for SAL-100

SEQ ID NO: 215 is the predicted amino acid sequence for SAL-105

SEQ ID NO: 216 is a second predicted amino acid sequence for SAL-50

25   DETAILED DESCRIPTION OF THE INVENTION

As noted above, the present invention is generally directed to compositions
and methods for the therapy of lung cancer. The compositions described herein include
polypeptides, fusion proteins and polynucleotides. Also included within the present invention
are molecules (such as an antibody or fragment thereof) that bind to the inventive

30   polypeptides. Such molecules are referred to herein as "binding agents."

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of the proteins described herein may be identified in antibody binding assays. Such assays may generally be performed using any of a variety of means known to those of ordinary skill in the art, as described, for example, in Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, 1988. For example, a polypeptide may be immobilized on a solid support (as described below) and contacted with patient sera to allow binding of antibodies within the sera to the immobilized polypeptide. Unbound sera may then be removed and bound antibodies detected using, for example, ^{125}I-labeled Protein A. Alternatively, a polypeptide may be used to generate monoclonal and polyclonal antibodies for use in detection of the polypeptide in blood or other fluids of lung cancer patients. Methods for preparing and identifying immunogenic portions of antigens of known sequence are well known in the art and include those summarized in Paul, *Fundamental Immunology*, 3rd ed., Raven Press, 1993, pp. 243-247.

The term "polynucleotide(s)," as used herein, means a single or double-stranded polymer of deoxyribonucleotide or ribonucleotide bases and includes DNA and corresponding RNA molecules, including HnRNA and mRNA molecules, both sense and anti-sense strands, and comprehends cDNA, genomic DNA and recombinant DNA, as well as wholly or partially synthesized polynucleotides. An HnRNA molecule contains introns and corresponds to a DNA molecule in a generally one-to-one manner. An mRNA molecule corresponds to an HnRNA and DNA molecule from which the introns have been excised. A polynucleotide may consist of an entire gene, or any portion thereof. Operable anti-sense polynucleotides may comprise a fragment of the corresponding polynucleotide, and the definition of "polynucleotide" therefore includes all such operable anti-sense fragments.

The compositions and methods of the present invention also encompass variants of the above polypeptides and polynucleotides.

A polypeptide "variant," as used herein, is a polypeptide that differs from the recited polypeptide only in conservative substitutions and/or modifications, such that the antigenic properties of the polypeptide are retained. In a preferred embodiment, variant polypeptides differ from an identified sequence by substitution, deletion or addition of five amino acids or fewer. Such variants may generally be identified by modifying one of the above polypeptide sequences, and evaluating the antigenic properties of the modified polypeptide using, for example, the representative procedures described herein. Polypeptide

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SSC, 0.5% SDS, 1.0 mM EDTA (pH 8.0); hybridizing at 50°C-65°C, 5X SSC, overnight or, in the event of cross-species homology, at 45°C with 0.5X SSC; followed by washing twice at 65°C for 20 minutes with each of 2X, 0.5X and 0.2X SSC containing 0.1% SDS. Such hybridizing DNA sequences are also within the scope of this invention, as are nucleotide

5   sequences that, due to code degeneracy, encode an immunogenic polypeptide that is encoded by a hybridizing DNA sequence.

Two nucleotide or polypeptide sequences are said to be "identical" if the sequence of nucleotides or amino acid residues in the two sequences is the same when aligned for maximum correspondence as described below. Comparisons between two sequences are

10   typically performed by comparing the sequences over a comparison window to identify and compare local regions of sequence similarity. A "comparison window" as used herein, refers to a segment of at least about 20 contiguous positions, usually 30 to about 75, 40 to about 50, in which a sequence may be compared to a reference sequence of the same number of contiguous positions after the two sequences are optimally aligned.

15   Optimal alignment of sequences for comparison may be conducted using the Megalign program in the Lasergene suite of bioinformatics software (DNASTAR, Inc., Madison, WI), using default parameters. This program embodies several alignment schemes described in the following references: Dayhoff, M.O. (1978) A model of evolutionary change in proteins – Matrices for detecting distant relationships. In Dayhoff, M.O. (ed.) Atlas of

20   Protein Sequence and Structure, National Biomedical Resarch Foundaiton, Washington DC Vol. 5, Suppl. 3, pp. 345-358; Hein J. (1990) Unified Approach to Alignment and Phylogenes pp. 626-645 *Methods in Enzymology* vol. 183, Academic Press, Inc., San Diego, CA; Higgins, D.G. and Sharp, P.M. (1989) Fast and sensitive multiple sequence alignments on a microcomputer *CABIOS* 5:151-153; Myers, E.W. and Muller W. (1988) Optimal alignments

25   in linear space *CABIOS* 4:11-17; Robinson, E.D. (1971) *Comb. Theor* 11:105; Santou, N. Nes, M. (1987) The neighbor joining method. A new method for reconstructing phylogenetic trees *Mol. Biol. Evol.* 4:406-425; Sneath, P.H.A. and Sokal, R.R. (1973) *Numerical Taxonomy – the Principles and Practice of Numerical Taxonomy*, Freeman Press, San Francisco, CA; Wilbur, W.J. and Lipman, D.J. (1983) Rapid similarity searches of nucleic

30   acid and protein data banks *Proc. Natl. Acad., Sci. USA* 80:726-730.

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libraries prepared from SCID mice with mouse anti-tumor sera, as described below in Example 4. Examples of cDNA sequences that may be isolated using this technique are provided in SEQ ID NO: 149-181.

5   A gene encoding a polypeptide described herein (or a portion thereof) may, alternatively, be amplified from human genomic DNA, or from lung tumor cDNA, via polymerase chain reaction. For this approach, sequence-specific primers may be designed based on the nucleotide sequences provided herein and may be purchased or synthesized. An amplified portion of a specific nucleotide sequence may then be used to isolate the full length gene from a human genomic DNA library or from a lung tumor cDNA library, using well 10 known techniques, such as those described in Sambrook et al., *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY (1989).

Once a DNA sequence encoding a polypeptide is obtained, the polypeptide may be produced recombinantly by inserting the DNA sequence into an expression vector and expressing the polypeptide in an appropriate host. Any of a variety of expression vectors 15 known to those of ordinary skill in the art may be employed to express recombinant polypeptides of this invention. Expression may be achieved in any appropriate host cell that has been transformed or transfected with an expression vector containing a polynucleotide that encodes the recombinant polypeptide. Suitable host cells include prokaryotes, yeast and higher eukaryotic cells. Preferably, the host cells employed are *E. coli*, yeast or a mammalian 20 cell line, such as COS or CHO cells. The DNA sequences expressed in this manner may encode naturally occurring polypeptides, portions of naturally occurring polypeptides, or other variants thereof. Supernatants from suitable host/vector systems which secrete the recombinant polypeptide may be first concentrated using a commercially available filter. The concentrate may then be applied to a suitable purification matrix, such as an affinity matrix or 25 ion exchange resin. Finally, one or more reverse phase HPLC steps can be employed to further purify the recombinant polypeptide.

Such techniques may also be used to prepare polypeptides comprising portions or variants of the native polypeptides. Portions and other variants having fewer than about 100 amino acids, and generally fewer than about 50 amino acids, may be generated using 30 techniques well known to those of ordinary skill in the art. For example, such polypeptides may be synthesized using any of the commercially available solid-phase techniques, such as

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extended conformation; (2) their inability to adopt a secondary structure that could interact with functional epitopes on the first and second polypeptides; and (3) the lack of hydrophobic or charged residues that might react with the polypeptide functional epitopes. Preferred peptide linker sequences contain Gly, Asn and Ser residues. Other near neutral amino acids, such as Thr and Ala may also be used in the linker sequence. Amino acid sequences which may be usefully employed as linkers include those disclosed in Maratea et al., *Gene* 40:39-46, 1985; Murphy et al., *Proc. Natl. Acad. Sci. USA* 83:8258-8262, 1986; U.S. Patent No. 4,935,233 and U.S. Patent No. 4,751,180. The linker sequence may be from 1 to about 50 amino acids in length. Peptide sequences are not required when the first and second polypeptides have non-essential N-terminal amino acid regions that can be used to separate the functional domains and prevent steric interference.

The ligated DNA sequences are operably linked to suitable transcriptional or translational regulatory elements. The regulatory elements responsible for expression of DNA are located only 5' to the DNA sequence encoding the first polypeptides. Similarly, stop codons require to end translation and transcription termination signals are only present 3' to the DNA sequence encoding the second polypeptide.

Fusion proteins are also provided that comprise a polypeptide of the present invention together with an unrelated immunogenic protein. Preferably the immunogenic protein is capable of eliciting a recall response. Examples of such proteins include tetanus, tuberculosis and hepatitis proteins (see, for example, Stoute et al. *New Engl. J. Med.*, 336:86-91 (1997)).

Polypeptides that comprise an immunogenic portion of a lung tumor protein may generally be used for therapy of lung cancer, wherein the polypeptide stimulates the patient's own immune response to lung tumor cells. The present invention thus provides methods for using one or more of the compounds described herein (which may be polypeptides, polynucleotides or fusion proteins) for immunotherapy of lung cancer in a patient. As used herein, a "patient" refers to any warm-blooded animal, preferably a human. A patient may be afflicted with disease, or may be free of detectable disease. Accordingly, the compounds disclosed herein may be used to treat lung cancer or to inhibit the development of lung cancer. In a preferred embodiment, the compounds are administered

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ordinary skill in the art. The DNA may also be "naked," as described, for example, in published PCT application WO 90/11092, and Ulmer et al., *Science* 259:1745-1749, 1993, reviewed by Cohen, *Science* 259:1691-1692, 1993. The uptake of naked DNA may be increased by coating the DNA onto biodegradable beads, which are efficiently transported

5   into the cells.

Routes and frequency of administration, as well as dosage, will vary from individual to individual and may parallel those currently being used in immunotherapy of other diseases. In general, the pharmaceutical compositions and vaccines may be administered by injection (*e.g.*, intracutaneous, intramuscular, intravenous or subcutaneous),

10   intranasally (*e.g.*, by aspiration) or orally. Between 1 and 10 doses may be administered over a 3-24 week period. Preferably, 4 doses are administered, at an interval of 3 months, and booster administrations may be given periodically thereafter. Alternate protocols may be appropriate for individual patients. A suitable dose is an amount of polypeptide or DNA that is effective to raise an immune response (cellular and/or humoral) against lung tumor cells in

15   a treated patient. A suitable immune response is at least 10-50% above the basal (*i.e.*, untreated) level. In general, the amount of polypeptide present in a dose (or produced *in situ* by the DNA in a dose) ranges from about 1 pg to about 100 mg per kg of host, typically from about 10 pg to about 1 mg, and preferably from about 100 pg to about 1 µg. Suitable dose sizes will vary with the size of the patient, but will typically range from about 0.01 mL to

20   about 5 mL.

While any suitable carrier known to those of ordinary skill in the art may be employed in the pharmaceutical compositions of this invention, the type of carrier will vary depending on the mode of administration. For parenteral administration, such as subcutaneous injection, the carrier preferably comprises water, saline, alcohol, a lipid, a wax

25   and/or a buffer. For oral administration, any of the above carriers or a solid carrier, such as mannitol, lactose, starch, magnesium stearate, sodium saccharine, talcum, cellulose, glucose, sucrose, and/or magnesium carbonate, may be employed. Biodegradable microspheres (*e.g.*, polylactic glycolide) may also be employed as carriers for the pharmaceutical compositions of this invention. Suitable biodegradable microspheres are disclosed, for example, in U.S.

30   Patent Nos. 4.897,268 and 5,075,109.

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(Natural Killer cells, lymphokine-activated killer cells), B cells, or antigen presenting cells (such as dendritic cells and macrophages) expressing the disclosed antigens. The polypeptides disclosed herein may also be used to generate antibodies or anti-idiotypic antibodies (as in U.S. Patent No. 4,918,164), for passive immunotherapy.

5   The predominant method of procuring adequate numbers of T-cells for adoptive immunotherapy is to grow immune T-cells in vitro. Culture conditions for expanding single antigen-specific T-cells to several billion in number with retention of antigen recognition in vivo are well known in the art. These in vitro culture conditions typically utilize intermittent stimulation with antigen, often in the presence of cytokines, such

10   as IL-2, and non-dividing feeder cells. As noted above, the immunoreactive polypeptides described herein may be used to rapidly expand antigen-specific T cell cultures in order to generate sufficient number of cells for immunotherapy. In particular, antigen-presenting cells, such as dendritic, macrophage or B-cells, may be pulsed with immunoreactive polypeptides or transfected with a polynucleotide sequence(s), using standard techniques well

15   known in the art. For cultured T-cells to be effective in therapy, the cultured T-cells must be able to grow and distribute widely and to survive long term in vivo. Studies have demonstrated that cultured T-cells can be induced to grow in vivo and to survive long term in substantial numbers by repeated stimulation with antigen supplemented with IL-2 (see, for example, Cheever et al. Ibid).

20   The polypeptides disclosed herein may also be employed to generate and/or isolate tumor-reactive T-cells, which can then be administered to the patient. In one technique, antigen-specific T-cell lines may be generated by in vivo immunization with short peptides corresponding to immunogenic portions of the disclosed polypeptides. The resulting antigen specific CD8+ CTL clones may be isolated from the patient, expanded using standard

25   tissue culture techniques, and returned to the patient.

Alternatively, peptides corresponding to immunogenic portions of the polypeptides may be employed to generate tumor reactive T cell subsets by selective in vitro stimulation and expansion of autologous T cells to provide antigen-specific T cells which may be subsequently transferred to the patient as described, for example, by Chang et al.

30   (Crit. Rev. Oncol. Hematol., 22(3), 213, 1996).

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at least about 80%, and preferably at least about 90%) of the patients for which lung cancer would be indicated using the full length protein, and that indicate the absence of lung cancer in substantially all of those samples that would be negative when tested with full length protein. The representative assays described below, such as the two-antibody sandwich assay, may generally be employed for evaluating the ability of a binding agent to detect metastatic human lung tumors.

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The ability of a polypeptide prepared as described herein to generate antibodies capable of detecting primary or metastatic human lung tumors may generally be evaluated by raising one or more antibodies against the polypeptide (using, for example, a representative method described herein) and determining the ability of such antibodies to detect such tumors in patients. This determination may be made by assaying biological samples from patients with and without primary or metastatic lung cancer for the presence of a polypeptide that binds to the generated antibodies. Such test assays may be performed, for example, using a representative procedure described below. Polypeptides that generate antibodies capable of detecting at least 20% of primary or metastatic lung tumors by such procedures are considered to be useful in assays for detecting primary or metastatic human lung tumors. Polypeptide specific antibodies may be used alone or in combination to improve sensitivity.

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Polypeptides capable of detecting primary or metastatic human lung tumors may be used as markers for diagnosing lung cancer or for monitoring disease progression in patients. In one embodiment, lung cancer in a patient may be diagnosed by evaluating a biological sample obtained from the patient for the level of one or more of the above polypeptides, relative to a predetermined cut-off value. As used herein, suitable "biological samples" include blood, sera, urine and/or lung secretions.

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The level of one or more of the above polypeptides may be evaluated using any binding agent specific for the polypeptide(s). A "binding agent," in the context of this invention, is any agent (such as a compound or a cell) that binds to a polypeptide as described above. As used herein, "binding" refers to a noncovalent association between two separate molecules (each of which may be free (i.e., in solution) or present on the surface of a cell or a solid support), such that a "complex" is formed. Such a complex may be free or immobilized (either covalently or noncovalently) on a support material. The ability to bind may generally

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be immobilized on the solid support using a variety of techniques known to those of skill in the art, which are amply described in the patent and scientific literature. In the context of the present invention, the term "immobilization" refers to both noncovalent association, such as adsorption, and covalent attachment (which may be a direct linkage between the antigen and
5   functional groups on the support or may be a linkage by way of a cross-linking agent). Immobilization by adsorption to a well in a microtiter plate or to a membrane is preferred. In such cases, adsorption may be achieved by contacting the binding agent, in a suitable buffer, with the solid support for a suitable amount of time. The contact time varies with temperature, but is typically between about 1 hour and about 1 day. In general, contacting a
10   well of a plastic microtiter plate (such as polystyrene or polyvinylchloride) with an amount of binding agent ranging from about 10 ng to about 10 μg, and preferably about 100 ng to about 1 μg, is sufficient to immobilize an adequate amount of binding agent.

Covalent attachment of binding agent to a solid support may generally be achieved by first reacting the support with a bifunctional reagent that will react with both the
15   support and a functional group, such as a hydroxyl or amino group, on the binding agent. For example, the binding agent may be covalently attached to supports having an appropriate polymer coating using benzoquinone or by condensation of an aldehyde group on the support with an amine and an active hydrogen on the binding partner (see, e.g., Pierce Immunotechnology Catalog and Handbook, 1991, at A12-A13).

20   In certain embodiments, the assay is a two-antibody sandwich assay. This assay may be performed by first contacting an antibody that has been immobilized on a solid support, commonly the well of a microtiter plate, with the sample, such that polypeptides within the sample are allowed to bind to the immobilized antibody. Unbound sample is then removed from the immobilized polypeptide-antibody complexes and a second antibody
25   (containing a reporter group) capable of binding to a different site on the polypeptide is added. The amount of second antibody that remains bound to the solid support is then determined using a method appropriate for the specific reporter group.

More specifically, once the antibody is immobilized on the support as described above, the remaining protein binding sites on the support are typically blocked.
30   Any suitable blocking agent known to those of ordinary skill in the art, such as bovine serum albumin or Tween 20™ (Sigma Chemical Co., St. Louis, MO). The immobilized antibody is

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that corresponds to a predetermined cut-off value. In one preferred embodiment, the cut-off value is the average mean signal obtained when the immobilized antibody is incubated with samples from patients without lung cancer. In general, a sample generating a signal that is three standard deviations above the predetermined cut-off value is considered positive for

5   lung cancer. In an alternate preferred embodiment, the cut-off value is determined using a Receiver Operator Curve, according to the method of Sackett et al., *Clinical Epidemiology: A Basic Science for Clinical Medicine*, Little Brown and Co., 1985, p. 106-7. Briefly, in this embodiment, the cut-off value may be determined from a plot of pairs of true positive rates (*i.e.*, sensitivity) and false positive rates (100%-specificity) that correspond to each possible

10   cut-off value for the diagnostic test result. The cut-off value on the plot that is the closest to the upper left-hand corner (*i.e.*, the value that encloses the largest area) is the most accurate cut-off value, and a sample generating a signal that is higher than the cut-off value determined by this method may be considered positive. Alternatively, the cut-off value may be shifted to the left along the plot, to minimize the false positive rate, or to the right, to

15   minimize the false negative rate. In general, a sample generating a signal that is higher than the cut-off value determined by this method is considered positive for lung cancer.

In a related embodiment, the assay is performed in a flow-through or strip test format, wherein the antibody is immobilized on a membrane, such as nitrocellulose. In the flow-through test, polypeptides within the sample bind to the immobilized antibody as the

20   sample passes through the membrane. A second, labeled antibody then binds to the antibody-polypeptide complex as a solution containing the second antibody flows through the membrane. The detection of bound second antibody may then be performed as described above. In the strip test format, one end of the membrane to which antibody is bound is immersed in a solution containing the sample. The sample migrates along the membrane

25   through a region containing second antibody and to the area of immobilized antibody. Concentration of second antibody at the area of immobilized antibody indicates the presence of lung cancer. Typically, the concentration of second antibody at that site generates a pattern, such as a line, that can be read visually. The absence of such a pattern indicates a negative result. In general, the amount of antibody immobilized on the membrane is selected

30   to generate a visually discernible pattern when the biological sample contains a level of polypeptide that would be sufficient to generate a positive signal in the two-antibody

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of immortal cell lines capable of producing antibodies having the desired specificity (*i.e.*, reactivity with the polypeptide of interest). Such cell lines may be produced, for example, from spleen cells obtained from an animal immunized as described above. The spleen cells are then immortalized by, for example, fusion with a myeloma cell fusion partner, preferably one that is syngeneic with the immunized animal. A variety of fusion techniques may be employed. For example, the spleen cells and myeloma cells may be combined with a nonionic detergent for a few minutes and then plated at low density on a selective medium that supports the growth of hybrid cells, but not myeloma cells. A preferred selection technique uses HAT (hypoxanthine, aminopterin, thymidine) selection. After a sufficient time, usually about 1 to 2 weeks, colonies of hybrids are observed. Single colonies are selected and tested for binding activity against the polypeptide. Hybridomas having high reactivity and specificity are preferred.

Monoclonal antibodies may be isolated from the supernatants of growing hybridoma colonies. In addition, various techniques may be employed to enhance the yield, such as injection of the hybridoma cell line into the peritoneal cavity of a suitable vertebrate host, such as a mouse. Monoclonal antibodies may then be harvested from the ascites fluid or the blood. Contaminants may be removed from the antibodies by conventional techniques, such as chromatography, gel filtration, precipitation, and extraction. The polypeptides of this invention may be used in the purification process in, for example, an affinity chromatography step.

Monoclonal antibodies of the present invention may also be used as therapeutic reagents, to diminish or eliminate lung tumors. The antibodies may be used on their own (for instance, to inhibit metastases) or coupled to one or more therapeutic agents. Suitable agents in this regard include radionuclides, differentiation inducers, drugs, toxins, and derivatives thereof. Preferred radionuclides include ^{90}Y, ^{123}I, ^{125}I, ^{131}I, ^{186}Re, ^{188}Re, ^{211}At, and ^{212}Bi. Preferred drugs include methotrexate, and pyrimidine and purine analogs. Preferred differentiation inducers include phorbol esters and butyric acid. Preferred toxins include ricin, abrin, diptheria toxin, cholera toxin, gelonin, Pseudomonas exotoxin, Shigella toxin, and pokeweed antiviral protein.

A therapeutic agent may be coupled (*e.g.*, covalently bonded) to a suitable monoclonal antibody either directly or indirectly (*e.g.*, via a linker group). A direct reaction

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be prepared in a variety of ways. For example, more than one agent may be coupled directly to an antibody molecule, or linkers which provide multiple sites for attachment can be used. Alternatively, a carrier can be used.

A carrier may bear the agents in a variety of ways, including covalent bonding
5   either directly or via a linker group. Suitable carriers include proteins such as albumins (e.g., U.S. Patent No. 4,507,234, to Kato et al.), peptides and polysaccharides such as aminodextran (e.g., U.S. Patent No. 4,699,784, to Shih et al.). A carrier may also bear an agent by noncovalent bonding or by encapsulation, such as within a liposome vesicle (e.g., U.S. Patent Nos. 4,429,008 and 4,873,088). Carriers specific for radionuclide agents include
10   radiohalogenated small molecules and chelating compounds. For example, U.S. Patent No. 4,735,792 discloses representative radiohalogenated small molecules and their synthesis. A radionuclide chelate may be formed from chelating compounds that include those containing nitrogen and sulfur atoms as the donor atoms for binding the metal, or metal oxide, radionuclide. For example, U.S. Patent No. 4,673,562, to Davison et al. discloses
15   representative chelating compounds and their synthesis.

A variety of routes of administration for the antibodies and immunoconjugates may be used. Typically, administration will be intravenous, intramuscular, subcutaneous or in the bed of a resected tumor. It will be evident that the precise dose of the antibody/immunoconjugate will vary depending upon the antibody used, the antigen density
20   on the tumor. and the rate of clearance of the antibody.

Diagnostic reagents of the present invention may also comprise DNA sequences encoding one or more of the above polypeptides, or one or more portions thereof. For example, at least two oligonucleotide primers may be employed in a polymerase chain reaction (PCR) based assay to amplify lung tumor-specific cDNA derived from a biological
25   sample, wherein at least one of the oligonucleotide primers is specific for a polynucleotide encoding a lung tumor protein of the present invention. The presence of the amplified cDNA is then detected using techniques well known in the art, such as gel electrophoresis. Similarly, oligonucleotide probes specific for a polynucleotide encoding a lung tumor protein of the present invention may be used in a hybridization assay to detect the presence of an inventive
30   polypeptide in a biological sample.

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The following Examples are offered by way of illustration and not by way of limitation.

<div align="center">EXAMPLES</div>

<div align="center">Example 1</div>

<div align="center">PREPARATION OF LUNG TUMOR-SPECIFIC cDNA SEQUENCES USING
DIFFERENTIAL DISPLAY RT-PCR</div>

This example illustrates the preparation of cDNA molecules encoding lung tumor-specific polypeptides using a differential display screen.

Tissue samples were prepared from breast tumor and normal tissue of a patient with lung cancer that was confirmed by pathology after removal of samples from the patient. Normal RNA and tumor RNA was extracted from the samples and mRNA was isolated and converted into cDNA using a $(dT)_{12}AG$ (SEQ ID NO: 47) anchored 3' primer. Differential display PCR was then executed using a randomly chosen primer (SEQ ID NO: 48). Amplification conditions were standard buffer containing 1.5 mM $MgCl_2$, 20 pmol of primer, 500 pmol dNTP and 1 unit of Taq DNA polymerase (Perkin-Elmer, Branchburg, NJ). Forty cycles of amplification were performed using 94 °C denaturation for 30 seconds, 42 °C annealing for 1 minute and 72 °C extension for 30 seconds. Bands that were repeatedly observed to be specific to the RNA fingerprint pattern of the tumor were cut out of a silver stained gel, subcloned into the pGEM-T vector (Promega, Madison, WI) and sequenced. The isolated 3' sequences are provided in SEQ ID NO: 1-16.

Comparison of these sequences to those in the public databases using the BLASTN program, revealed no significant homologies to the sequences provided in SEQ ID NO: 1-11. To the best of the inventors' knowledge, none of the isolated DNA sequences have previously been shown to be expressed at a greater level in human lung tumor tissue than in normal lung tissue.

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aminopeptidase. Clone LT86-9 appears to contain two inserts, with the 5' sequence showing homology to the previously identified antisense sequence of interferon alpha-induced P27, and the 3' sequence being similar to LT86-6. Clone LT86-14 (SEQ ID NO: 30) was found to show some homology to the trithorax gene and has an "RGD" cell attachment sequence and a

5   beta-Lactamase A site which functions in hydrolysis of penicillin. Clones LT86-1, LT86-2, LT86-4, LT86-5 and LT86-10 (SEQ ID NOS: 17, 18; 20, 21 and 26, respectively) were found to show homology to previously identified genes. A subsequently determined extended cDNA sequence for LT86-4 is provided in SEQ ID NO: 66, with the corresponding predicted amino acid sequence being provided in SEQ ID NO: 67.

10   Subsequent studies led to the isolation of five additional clones, referred to as LT86-20, LT86-21, LT86-22, LT86-26 and LT86-27. The determined 5' cDNA sequences for LT86-20, LT86-22, LT86-26 and LT86-27 are provided in SEQ ID NO: 68 and 70-72, respectively, with the determined 3' cDNA sequences for LT86-21 being provided in SEQ ID NO: 69. The corresponding predicted amino acid sequences for LT86-20, LT86-21, LT86-

15   22, LT86-26 and LT86-27 are provided in SEQ ID NO: 73-77, respectively. LT86-22 and LT86-27 were found to be highly similar to each other. Comparison of these sequences to those in the gene bank as described above, revealed no significant homologies to LT86-22 and LT86-27. LT86-20, LT86-21 and LT86-26 were found to show homology to previously identified genes.

20

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predicted amino acid sequences are provided in SEQ ID NO: 93-101, respectively. L86S-30, L86S-39 and L86S-47 were found to be similar to each other. Comparison of these sequences with those in the gene bank as described above, revealed no significant homologies to L86S-14. L86S-29 was found to show some homology to a previously identified EST.

5   L86S-6, L86S-11, L86S-34, L86S-39, L86S-47, L86S-49 and L86S-51 were found to show some homology to previously identified genes.

In further studies, a directional cDNA library was constructed using a Stratagene kit with a Lambda Zap Express vector. Total RNA for the library was isolated from two primary squamous lung tumors and poly A+ RNA was isolated using an oligo dT

10   column. Antiserum was developed in normal mice using a pool of sera from three SCID mice implanted with human squamous lung carcinomas. Approximately 700,000 PFUs were screened from the unamplified library with *E. coli* absorbed mouse anti-SCID tumor serum. Positive plaques were identified as described above. Phage was purified and phagemid excised for 180 clones with inserts in a pBK-CMV vector for expression in prokaryotic or

15   eukaryotic cells.

The determined cDNA sequences for 23 of the isolated clones are provided in SEQ ID NO: 126-148. Comparison of these sequences with those in the public database as described above revealed no significant homologies to the sequences of SEQ ID NO: 139 and 143-148. The sequences of SEQ ID NO: 126-138 and 140-142 were found to show

20   homology previously identified human polynucleotide sequences.

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tags (ESTs). The sequences of SEQ ID NO: 150, 155 and 159-181 were found to show homology to sequences previously identified in humans.

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Example 6

ISOLATION OF DNA SEQUENCES ENCODING LUNG TUMOR ANTIGENS

DNA sequences encoding antigens potentially involved in squamous cell lung
5   tumor formation were isolated as follows.

A lung tumor directional cDNA expression library was constructed employing
the Lambda ZAP Express expression system (Stratagene, La Jolla, CA). Total RNA for the
library was taken from a pool of two human squamous epithelial lung carcinomas and poly
A+ RNA was isolated using oligo-dT cellulose (Gibco BRL, Gaithersburg, MD). Phagemid
10   were rescued at random and the cDNA sequences of isolated clones were determined.

The determined cDNA sequence for the clone SLT-T1 is provided in SEQ ID
NO: 102, with the determined 5' cDNA sequences for the clones SLT-T2, SLT-T3, SLT-T5,
SLT-T7, SLT-T9, SLT-T10, SLT-T11 and SLT-T12 being provided in SEQ ID NO: 103-
110, respectively. The corresponding predicted amino acid sequence for SLT-T1, SLT-T2,
15   SLT-T3, SLT-T10 and SLT-T12 are provided in SEQ ID NO: 111-115, respectively.
Comparison of the sequences for SLT-T2, SLT-T3, SLT-T5, SLT-T7, SLT-T9 and SLT-T11
with those in the public databases as described above, revealed no significant homologies.
The sequences for SLT-T10 and SLT-T12 were found to show some homology to sequences
previously identified in humans.

20   The sequence of SLT-T1 was determined to show some homology to a PAC
clone of unknown protein function. The cDNA sequence of SLT-T1 (SEQ ID NO: 102) was
found to contain a mutator (MUTT) domain. Such domains are known to function in removal
of damaged guanine from DNA that can cause A to G transversions (see, for example, el-
Deiry, W.S., 1997 *Curr. Opin. Oncol.* 9:79-87; Okamoto, K. et al. 1996 *Int. J. Cancer*
25   65:437-41; Wu, C. et al. 1995 *Biochem. Biophys. Res. Commun.* 214:1239-45; Porter, D.W.
et al. 1996 *Chem. Res. Toxicol.* 9:1375-81). SLT-T1 may thus be of use in the treatment, by
gene therapy, of lung cancers caused by, or associated with, a disruption in DNA repair.

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Example 7

SYNTHESIS OF POLYPEPTIDES

Polypeptides may be synthesized on a Perkin Elmer/Applied Biosystems
5   Division 430A peptide synthesizer using FMOC chemistry with HPTU (O-Benzotriazole-
N,N,N',N'-tetramethyluronium hexafluorophosphate) activation.   A Gly-Cys-Gly sequence
may be attached to the amino terminus of the peptide to provide a method of conjugation,
binding to an immobilized surface, or labeling of the peptide. Cleavage of the peptides from
the solid support may be carried out using the following cleavage mixture: trifluoroacetic
10   acid:ethanedithiol:thioanisole:water:phenol (40:1:2:2:3).   After cleaving for 2 hours, the
peptides may be precipitated in cold methyl-t-butyl-ether. The peptide pellets may then be
dissolved in water containing 0.1% trifluoroacetic acid (TFA) and lyophilized prior to
purification by C18 reverse phase HPLC.   A gradient of 0%-60% acetonitrile (containing
0.1% TFA) in water (containing 0.1% TFA) may be used to elute the peptides. Following
15   lyophilization of the pure fractions, the peptides may be characterized using electrospray or
other types of mass spectrometry and by amino acid analysis.

From the foregoing, it will be appreciated that, although specific embodiments
of the invention have been described herein for the purposes of illustration, various
20   modifications may be made without deviating from the spirit and scope of the invention.

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9.   A vaccine comprising the polypeptide of claim 2 and an immune response enhancer.

10.   The vaccine of claim 9 wherein the immune response enhancer is an adjuvant.

11.   A vaccine comprising the polynucleotide of claims 1 or 4 and an immune response enhancer.

12.   The vaccine of claim 11 wherein the immune response enhancer is an adjuvant.

13.   A pharmaceutical composition for the treatment of lung cancer comprising a polypeptide and a physiologically acceptable carrier, the polypeptide comprising an immunogenic portion of a lung protein or of a variant thereof, wherein said protein comprises an amino acid sequence encoded by a polynucleotide comprising a sequence selected from the group consisting of:

(a)   sequences recited in SEQ ID NO: 12-18, 20, 21, 26, 49, 50, 52, 54, 64, 66, 68, 69, 71, 78, 84, 85, 88, 91, 92, 116-120, 126-138, 140-142, 150, 155 and 159-181;

(b)   sequences complementary to the sequences of SEQ ID NO: 12-18, 20, 21, 26, 49, 50, 52, 54, 64, 66, 68, 69, 71, 78, 84, 85, 88, 91, 92, 116-120, 126-138, 140-142, 150, 155 and 159-181; and

(c)   variants of the sequences of (a) and (b).

14.   A vaccine for the treatment of lung cancer comprising a polypeptide and an immune response enhancer, said polypeptide comprising an immunogenic portion of a lung protein or of a variant thereof, wherein said protein comprises an amino acid sequence encoded by a polynucleotide comprising a sequence selected from the group consisting of:

(a)   sequences recited in SEQ ID NO: 12-18, 20, 21, 26, 49, 50, 52, 54, 64, 66, 68, 69, 71, 78, 84, 85, 88, 91, 92, 116-120, 126-138, 140-142, 150, 155 and 159-181;

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21.   A pharmaceutical composition comprising a fusion protein according
to any one of claims 18-20 and a physiologically acceptable carrier.

5   22.   A vaccine comprising a fusion protein according to any one of claims
18-20 and an immune response enhancer.

23.   The vaccine of claim 22 wherein the immune response enhancer is
an adjuvant.

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24.   A method for inhibiting the development of lung cancer in a patient,
comprising administering to the patient an effective amount of the pharmaceutical
composition of claim 21.

15   25.   A method for inhibiting the development of lung cancer in a patient,
comprising administering to the patient an effective amount of the vaccine of claim 22.

26.   A method for inhibiting the development of lung cancer in a patient,
comprising administering to the patient a polynucleotide under conditions such that the
20   polynucleotide enters a cell of the patient and is expressed therein, the polynucleotide having
a sequence selected from the group consisting of:
        (a)   a sequence provided in SEQ ID NO: 102;
        (b)   sequences complementary to a sequence of SEQ ID NO: 102; and
        (c)   variants of the sequence of SEQ ID NO: 102.

25   27.   A method for detecting lung cancer in a patient, comprising:
        (a)   contacting a biological sample obtained from the patient with a binding
agent which is capable of binding to a polypeptide, the polypeptide comprising an
immunogenic portion of a lung tumor protein or a variant thereof, wherein said protein
comprises an amino acid sequence encoded by a polynucleotide comprising a nucleotide
30   sequence selected from the group consisting of sequences provided in SEQ ID NO: 1-31, 49-

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(a)   sequences recited in SEQ ID NO: 1-11, 19, 22-25, 27-31, 51, 53, 55, 63, 70, 72, 79, 80, 86, 87, 89, 90, 102-107, 109, 139, 143-149, 151-154 and 156-158;

(b)   the complements of nucleotide sequences recited in SEQ ID NO: 1-11, 19, 22-25, 27-31, 51, 53, 55, 63, 70, 72, 79, 80, 86, 87, 89, 90, 102-107, 109, 139, 143-149, 151-154 and 156-158; and

(c)   variants of the sequences of (a) and (b).

32.   A method for inhibiting the development of lung cancer in a patient, comprising administering to the patient a therapeutically effective amount of a monoclonal antibody according to claim 31.

33.   The method of claim 32 wherein the monoclonal antibody is conjugated to a therapeutic agent.

34.   A method for detecting lung cancer in a patient comprising:

(a)   obtaining a biological sample from the patient;

(b)   contacting the sample with at least two oligonucleotide primers in a polymerase chain reaction, wherein at least one of the oligonucleotides is specific for a polynucleotide encoding a polypeptide comprising an immunogenic portion of a lung tumor protein or a variant thereof, said protein comprising an amino acid sequence encoded by a polynucleotide comprising a nucleotide sequence selected from the group consisting of sequences recited in SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181, the complements of said sequences and variants thereof; and

(c)   detecting in the sample a DNA sequence that amplifies in the presence of the oligonucleotide primers, thereby detecting lung cancer.

35.   The method of claim 34, wherein at least one of the oligonucleotide primers comprises at least about 10 contiguous nucleotides of a polynucleotide comprising a sequence selected from SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181.

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provided in SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181, the complements of said sequences and variants thereof.

44.   A method for detecting lung cancer in a patient, comprising:

(a)   obtaining a biological sample from the patient;

5   (b)   contacting the biological sample with an oligonucleotide probe specific for a polynucleotide encoding a polypeptide comprising an immunogenic portion of a lung tumor protein or a variant thereof, said protein comprising an amino acid sequence encoded by a polynucleotide comprising a nucleotide sequence selected from the group consisting of sequences recited in SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-

10   120 and 126-181, the complements of said nucleotide sequences and variants thereof; and

(c)   detecting in the sample a DNA sequence that hybridizes to the oligonucleotide probe, thereby detecting lung cancer in the patient.

45.   The method of claim 44 wherein the oligonucleotide probe comprises at least about 15 contiguous nucleotides of a polynucleotide having a nucleotide sequence

15   selected from the group consisting of sequences recited in SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181, the complements of said nucleotide sequences and variants thereof.

46.   A diagnostic kit comprising an oligonucleotide probe specific for a polynucleotide encoding a polypeptide comprising an immunogenic portion of a lung tumor

20   protein or a variant thereof, said protein comprising an amino acid sequence encoded by a polynucleotide comprising a nucleotide sequence selected from the group consisting of sequences recited in SEQ ID NO: 1-31, 49-55, 63, 64, 66, 68-72, 78-80, 84-92, 102-110, 116-120 and 126-181, the complements of said sequences and variants thereof.

47.   The diagnostic kit of claim 46, wherein the oligonucleotide probe

25   comprises at least about 15 contiguous nucleotides of a polynucleotide having a nucleotide sequence selected from the group consisting of sequences recited in SEQ ID NO: 1-31, 49-55,

55

pharmaceutically acceptable carrier.

55.   A composition for the treatment of lung cancer in a patient, comprising
T cells proliferated in the presence of a polynucleotide of claim 1, in combination with a
pharmaceutically acceptable carrier.

56.   A method for treating lung cancer in a patient, comprising the steps of:
(a)   incubating antigen presenting cells in the presence of at least one
polypeptide of claim 2; and
(b)   administering to the patient the incubated antigen presenting cells.

57.   A method for treating lung cancer in a patient, comprising the steps of:
(a)   incubating antigen presenting cells in the presence of at least one
polynucleotide of claim 1; and
(b)   administering to the patient the incubated antigen presenting cells.

58.   The method of claims 54 or 55 wherein the antigen presenting cells are
selected from the group consisting of dendritic cells and macrophage cells.

59.   A composition for the treatment of lung cancer in a patient, comprising
antigen presenting cells incubated in the presence of a polypeptide of claim 2, in combination
with a pharmaceutically acceptable carrier.

60.   A composition for the treatment of lung cancer in a patient, comprising
antigen presenting cells incubated in the presence of a polynucleotide of claim 1, in
combination with a pharmaceutically acceptable carrier.

                                            1

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cgacggagaa nmagntctgt tattccatgg gaaattkgaa tttccttcyt tkaaatatct 480
taaaatnggt cctcctwaaa aaagttcggc tggggcccgk tggctcacat tttkttaycc 540
cyccccttt tggggargcc caarggccgg kttgawtnnc ccttgagggg ccanaactcc 600
agnaaccrgn cccgggccar smgwkgkstr armcccttc cyyccmaraa aawwcsmaaa 660
wwttycccsc cygsykggct ggkasckgtt myyyyygmtm csyagcttgc tt        712
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<210> 5
<211> 679
<212> DNA
<213> Homo sapiens

<400> 5
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acatccctt gtgagcacgg ctgctccgga atactgacca tctgggctag cacgacctaa 120
cagagggttc tgcaggatgt gctattttaa agcagctggg tgcaacttgt gaaaacggga 180
atctngaagc agaacatgtn atcagcgatg gctgggattg gtggacagga ttgacaggag 240
tatttgaggc tctaccaggc ctgtctacag gacagcttca tcgaagggac atttttttaac 300
ctgttatttt anatnccaca tatntttttt aatgctnaag catacaggtt gaatttctgg 360
atcgtaacta ctagtgactt ctgaggttta cagttngaat atgttctcnn aggttatca 420
agttntgtta ttgatgatng gtaatctaca cctctggaag ctgtngaatg tgaaaaagat 480
ncntncanct gaccagtttg nagggcactc tcttctggna agnaatccgn ccaaaaaaat 540
tgtttcnagg gggcntgggg ggttaaaaa aatgtttctn ttnccntaaa aatgtttacc 600
cnnctattga aaaaatgggg gtcgnggggg gcttnaaatc cccnanttnt gaatnttnta 660
tccggaanct tggtttccc                                            679
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<210> 6
<211> 369
<212> DNA
<213> Homo sapiens

<400> 6
tcagtccagt catgggtcct ataagagaag tcactctgtg agtttccatg gaggaagaaa 60
aagcttcatt tctttaccct gcagcaacag cggagggagg gagagcctat cttcctttgca 120
aattcattaa ctttgtggtt gaagggagca gcgtcngaaa ctgctttagc acagtgggag 180
gaaaacaaac agattcatct ccggaaacca aaggaaaggg tragtgggtt tttattagcc 240
agctgtatcc tagatggtca atttccagtg gatgaataca ccttacgtac gtttctcttg 300
cttcctacct nggcctgatc agctnggcac ttraatcatt ccgtnggggt wgctgtnaca 360
ctggactga 369

<210> 7
<211> 264
<212> DNA
<213> Homo sapiens

<400> 7
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aggaaaggat tacaggcgtg agccactgcg cccggcctct tctccacttt cataggttcc 120
agtctctggt tcttctttct cagtttgttg tttttgcttc ttaaatmatg gagatnagaa 180
tgaacactac actcggaatc aggaagccct gcctggcgcc tctgtcacct gtctaggggc 240
ttcttctcac tgagtcatcc agca 264

<210> 8
<211> 280
<212> DNA
<213> Homo sapiens

<400> 8
acctcaactg cccanaacan aactgttgta caagatttga ggatttaaca atatttcaca 60
tgaaatattt cagacctacg ngagggctta aagacnaatt aaatgagcac cngtgtgccc 120
accgccccna ttaagaatta gagcaagcag tgaggtgaag ccttgtcctt gcttttaaca 180
tagaaagtga tccaaattca ccaaacttga cttnnggttt tgcagtgtgg cctcctgatt 240
ctagacnctg gcgaaacatt tgatgggcaa aaaaaaaaaa 280

<210> 9
<211> 449
<212> DNA
<213> Homo sapiens

<400> 9
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ntgcagtgct natgactggc tttgcagttn attttgattc aggcaacaga tgttcctttt 120
ggttccctgt ctcccatggg cgtcatttca tgttgtcctc tgccttcccc cagatattct 180
aagttcagga cacaagcttc tggcccatgc agagcagagg ccatgagggg tcacagcatg 240
ggtacgggag gaaacactgg gctnacccag atnctggact tgagtcttgc ctctgctgct 300
tgctgcacag cttctgtcat ggtgctaaac ctgtgacctg cctcacaggc ttagagcatg 360
cccgtagaag tactctnaac taaratgctt tccacaaatg agatggtttc atgaaaactt 420
caaatagagg gcctgggcaa aaaaaaaaa 449

<210> 10
<211> 538
<212> DNA
<213> Homo sapiens

<400> 10
ttttttttttt ttcccaaagg cctcaraaca ctagtcttct aattccaagc agaaagttac 60

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atccgccggg atacatgcca cttggtttga taaatcaaaa tacagcatcc ttcagatccc 120
tttgctgagc aatacaatta tttgtatatg ttactttttt ttctgtttgg ctnaaagatt 180
tgatatgagc tgaggaaaat gaagccntta ctgctatnag atctnatccc tttccaccac 240
ctttcaggga tnttggcact gcayatattc agaattcccc nnagtcgctn gtgataaaaa 300
tgtcttcaga gatggcagaa tatgtttcct ttggtacatg ttcattaaaa atatacacgt 360
gctcactact gtggatatgt atgtnttgac cgatnacaca ggctgattta gggaagagat 420
aaaagcacac ttngaattta ttagcctttc accnagacta anattctgaa attaagaatg 480
tattccttgg tcaacaattt tcctcttctc ttagccctct tacattgtan tggactga   538
```

`<210> 11`
`<211> 543`
`<212> DNA`
`<213> Homo sapiens`

`<400> 11`
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caaccctcgc catcccagca aatcccctct ctcccttctc atgggagtgc cttgtattca 120
tcaggcatct gggacttgat gtgggtntgg gatttgaaat cagagcacct nggtctctst 180
caccattctn tcacttatta gctctnacct tgggtnaata cctgccttag tgtcntaggt 240
acaatatgaa tattgtctat ttctcaggga ttgcaatgac nagtnnatna gtgcatgaga 300
gggtaaaacc acagggtact ccgctcctcc naagaatgga gaattttttc tagaagccca 360
natntgcttg gaaggttggc caccnagagc cnnaatcttc ttttatttnc cactgaangc 420
ctaagaggna attctgaact catccccnna tgacctctcc cgaatmagaa tatctctggc 480
acttaccata ttttcttgcc ctcttccact tacnaaactc ctttattcct taacnggacg 540
aaa                                                             543
```

`<210> 12`
`<211> 329`
`<212> DNA`
`<213> Homo sapiens`

`<400> 12`
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cgatgacttg ggcagtgagt gggcctcctg ccaggtggca ggcacagct tagaccaaac 60
ccttggcctc ccccctctgc agstacctct gaccaagaag gaaactagca agcctatgct 120
ggcaagacca taggtggggt gctgggaatc ctcggggccg gctggcaccc actcctggtg 180
ctcaagggag agacccactt gttcagatgc atrggcctca ggcggttcaa ggcrgtctta 240
gagccacaga gtcaaataaa aatcaatttt gagagaccac agcacctgct gctttgatcg 300
tgatgttcaa ggcaagttgc aagtcatcg                                 329
```

`<210> 13`
`<211> 314`
`<212> DNA`
`<213> Homo sapiens`

`<400> 13`
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cgatgacttg cacccgggag ctgtgacagt ggcctggaag cagatggcag ccccgtcaag 60
gcgggagtgg agaccaccaa accctccaaa cagagcaaca actagtacgc ggccagcagc 120
tacctgagcc tgacgcccga gcagtggaag tcccacagaa gctacagctg ccaggtcacg 180
catgaaggga gcaccgtgga gaagacagtg gcccctacag aatgttcata ggttcccnac 240
tctnaccccca cccacgggag cctgganctg cangatcccg ggggaagggt ctctctcccc 300
atcccaagtc atcg                                                 314
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`<210> 14`
`<211> 691`
`<212> DNA`

<213> Homo sapiens

<400> 14
cgattacttg cacaatgcan attagaacccc aaatgaaggg tacaacccag atcttctggc 60
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gaagagaaga ataaagtcta ttttggtctt tggtagcchg ggtaangaga atgctstcac 180
tctacnagaa aaccccaagt gaacccggct aatcaggacc gtgcttggga agggagcagg 240
ggcattacct ttcaacacca gaggttcttt gccttctctc tgcagggact cgargactat 300
gtgaagtggc tgggargggca tcactcggct tggttcattg gtrttctcat cataaactat 360
natttctttg gaaaaagatc ctcttgaaag artccttgcc ttccctacag gaaatcaagt 420
ctaggacagt gatcttgccc ctgcttgcas tctccgccgg ctgatcttat csgscccagt 480
tkatgtgsam cgctccttgg atrtkactct tgttttwctc cvaggaaggg gcytgcmagt 540
ccnwtnaatg amssgggccc ttaactccgg scrggtnamy ncttgsctsc rattttgggt 600
ycytcttcyt ttgsccmggt tcktcnaaac cacttngttr aattccccgg sccgcctkgc 660
nggtycaacc wttttgggaa mamcyccccc c   691

<210> 15
<211> 355
<212> DNA
<213> Homo sapiens

<400> 15
acctgaactg tgtgttgaag agtgatgtcc tgctgcctgg agctcaagtc actactgatg 60
accgtgccta tgtccgacag ctagttncct ccatggatgt gactgagacc aatgtcttct 120
tcyaccctcg gctcttacct ttgacnaagt ctcccgttga gagtactacc gaaccaccag 180
cagttcgagc ctctnaagag cgtctaagcg atggggatat atatttactg gagaatgggc 240
tcaacctctt cctctgggtg ggagcaagcg tccagcaggg tgttgtccag agcctttttca 300
gcgtctcctc cttcagtcag atcaccagtg gtntgagtgt tctgccagtt caggt   355

<210> 16
<211> 522
<212> DNA
<213> Homo sapiens

<400> 16
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tgagagacag cgcatgctgg tgcagcgtan ggacgaactc ctccagcaag ctcgcagacg 120
tttcttgaac aaaagttctg aagatgatgc ggcctcagag agcttcctcc cctcggaagg 180
tgcgtcctct gaccccgtga ccctncgtcg aangatgctg gctgccgccg cggaacggan 240
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tgtgcccggc tgactggagg aggcctgtcc aattctcccc gccccatgga aaagcgggct 360
tgactgcatt gccgctgtat naaagcatgt ggtcttacag tgttnggacn gctnatnaat 420
ttnatcctnc tntgtaatac ttcctatgtg acatttctct tccccttgga aacactgcan 480
attttaactg tgagtttgat ctcttctngt gttactggac tg   522

<210> 17
<211> 317
<212> DNA
<213> Homo sapiens

<400> 17
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aaggataagc accagagaaa gaaggttcag ccggccgtcc tgaaatatta taaggtggat 120
gagaatggca aaattagttg ccttcgtcga gagtgcccct ctgatgaatg tggtgctggg 180
gtgtttatgg caagtcactt tgacagacat tattgtggca aatgttgtct gacccactgt 240

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ttcaactaac cagaagacaa gtaactgtat gagttaatta aagacatgaa ctaaaaaaaa 300
aaaaaaaaaa actcgag                                                   317
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<210> 18
<211> 392
<212> DNA
<213> Homo sapiens

<400> 18
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tggagatttc taatgaggtg aggaagttcc gtacattgac agaattgatc ctcgatgctc 60
aggaacatgt taaaaatcct tacaaaggca aaaaactcaa gaaacaccca gacttcccca 120
agaagcccct gacccctat ttccgcttct tcatggagaa gcgggccaag tatgcgaaac 180
tccaccctca gatgagcaac ctggacctga ccaagattct gtccaagaaa tacaaggagc 240
ttccggagaa gaagaagatg aaatatgttc cggacttcca gagaagagaa acaggagttc 300
gagcgaaacc tggcccgatt cagggaggat caccccccacc ttatccagaa tgccaagaat 360
cggacatccc agagaagccc caagaccccc cg              392
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<210> 19
<211> 2624
<212> DNA
<213> Homo sapiens

<400> 19
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gaaacagtga gaaggagatt cctgtgctca atgagctgcc agtccccatg gtggccgct 60
acattcgcat aaaccctcag tcctggtttg ataacgggag catctgcatg aggatggaga 120
tcttgggctg cccactgccg gatcctaata actattatca ccgacgtaat gagatgacca 180
ccacggatga cctggatttt aagcaccaca actattagga aatgcgccag ttgatgaagg 240
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gcctgaaatt gtatgcggta gagatctctg accatcctgg ggaacatgaa gttggtgagc 360
ccgagttcca ctacatcgca ggggcccacg gcaatgaggt tctgggacga gaactgctgc 420
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tcgatatcaa caacaacttt ccggatttaa actcgctgct ctgggaggca gaggaccagc 660
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gtctgagatt ctaaaaaggg tgcttgacca ctggccagga agggaaatca ggccttcccc 1860
catttgcgtg acattcaagc ttcccagtgc atttgcaagt ggcacagttg acattgcagc 1920
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<210> 20
<211> 488
<212> DNA
<213> Homo sapiens

<400> 20
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gctccgcagg gtgaggtggc tttgaccccg ggttgcccgg ccagcacgac cgaggaggtg 120
gctggacagc tggaggatga acggagaagc cgactgcccc acagacctgg aaatggccgc 180
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gaacaacctt ccatccaatg acagctccca gttcaaaacc acccaaacac acatggaccg 300
ggaaaaagtt gcattgaaag actttctgg agacatgtgc aagctcaaat gggtcgagat 360
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tgtttaaaat ccttacaaag gcaaaaaatc aagaaacacc ccgacttccc cgagaaagcc 480
cctaaccc                                                  488
```

<210> 21
<211> 391
<212> DNA
<213> Homo sapiens

<400> 21
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tggcgattgc tgatgctttg ggcaaaatcc ctcagacagt cctgtggcgg tacactggaa 120
cccgaccatc gaatcttgcg aacaacacga tacttgttca gtggctaccc caaaacgatc 180
tgcttggtca cccaatgacc cgtgcctttta tcacccatgc tagttcccat ggtgttaatg 240
aaagcatatg caatggcgtt cccatggtga tgatacccctt atttggtgat cagatggaca 300
atgcaaagcg caggagact aagggagctg gagtgaccct gaatgttctg gagatgactt 360
ctgaagatct agaagatgct ctgaagagca g                        391
```

<210> 22
<211> 1320
<212> DNA
<213> Homo sapiens

<400> 22
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gtcctggatt acttttcaga aagaagtaat cctttttatg acagaacatg taataatgaa 120
gtggtcaaaa tgcagaggct aacattagaa cacttgaatc agatggttgg aatcgagtac 180
atccttttgc atgctcaaga gcccattctt ttcatcattc ggaagcaaca gcggcagtcc 240
cctgcccaag ttatcccact agctgattac tatatcattg ctggagtgat ctatcaggca 300
ccagacttgg gatcagttat aaactctaga gtgcttactg cagtgcatgg tattcagtca 360
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gcttttgatg aagctatgtc atactgtcga tatcatcctt ccaaagggta ttggtggcac 420
ttcaaagatc atgaagagca agataaagtc agacctaaag ccaaaaggaa agaagaacca 480
agctctattt ttcagagaca acgtgtggat gctttacttt tagacctcag acaaaaattt 540
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aaagaggcag aacctatacc agaaactgta aaacctgagg agaaggagac cacaaagaat 660
gtacaacaga cagtgagtgc taaaggcccc cctgaaaaac ggatgagact tcagtgagta 720
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gtaacctttc ctttcccgga cttgagcaac ctacacactc acatgtttaa tggtagatat 1260
gttttaaagc aagataaagg tattggtttt aaaaaaaaaa aaaaaaaaaa aaaactcgag 1320
```

<210> 23
<211> 633
<212> DNA
<213> Homo sapiens

<400> 23
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tgatgggggt taagccgggg gaggacgcat cggggcctgc tgaagacctt gtgagaagat 120
ctgagaaaga tactgcagct gttgtctcca gacagggcag ctccctgaac ctctttgaag 180
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caatgaatgc cacggccacc aaggttgcta actgcagctt gggaactgcc accatcatcg 360
gtgagaactt gaacaatgag gtcatgatga agaaatacag cccctcggac cctgcatttg 420
catatgcgca gctgacccac gatgagctga ttcagctggt cctcaaacag aaggaaacga 480
taagcaagaa ggagttccag gtccgcgagc tggaagacta cattgacaac ctgctcgtca 540
gggtcatgga agaaaccccc aatatcctcc gcatcccgac tcaggttggc aaaaaagcag 600
gaaagatgta aattagcaga aaaaaaactc gag                              633
```

<210> 24
<211> 1328
<212> DNA
<213> Homo sapiens

<400> 24
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tgggttgaca gctcttggat ccctattttg aacagtggta gtgtcctgga ttacttttca 120
gaaagaagta atccttttta tgacagaaca tgtaataatg aagtggtcaa aatgcagagg 180
ctaacattag aacacttgaa tcagatggtt ggaatcgagt acatcctttt gcatgctcaa 240
gagcccattc tttttcatcat tcggaagcaa cagcggcagt cccctgccca agttatccca 300
ctagctgatt actatatcat tgctggagtg atctatcagg caccagactt gggatcagtt 360
ataaactcta gagtgcttac tgcagtgcat ggtattcagt cagcttttga tgaagctatg 420
tcatactgtc gatatcatcc ttccaaaggg tattggtggc acttcaaaga tcatgaagag 480
caagataaag tcagacctaa agccaaaagg aaagaagaac caagctctat ttttcagaga 540
caacgtgtgg atgctttact tttagacctc agacaaaaaa tttccaccca aatttgtgca 600
gtggatcaaa caaagaaaga ggcagaacct ataccagaaa ctgtaaaacc tgaggagaag 660
gagaccacaa agaatgtaca acagacagtg agtgctaaag gcccccctga aaaacggatg 720
agacttcagt gagtactgga caaaagagaa gcctggaaga ctcctcatgc tagttatcat 780
acctcagtac tgtggctctt gagctttgaa gtactttatt gtaaccttct tatttgtatg 840
```

```
gaatgcgctt atttttttga aaggatatta ggccggatgt ggtggctcac gcctgtaatc 900
ccagcacttt gggaggccat ggcgggtgga tcacttgagg tcagaagttc aagaccagcc 960
tgaccaatat ggtgaaaccc cgtctctact aaaaatacaa aaattagccg ggcgtggtgg 1020
cgggcgccca tagtcccagc tactcgggag gctgagacag gagacttgct tgaacccggg 1080
aggtggaggt tgccctgagc tgattatcat gctgttcac tccagcttgg gcgacagagc 1140
gagactttgt ctcaaaaaag aagaaaagat attattccca tcatgatttc ttgtgaatat 1200
ttgtgatatg tcttctgtaa ccttttcctct cccggacttg agcaacctac acactcacat 1260
gtttactggt agatatgttt aaaagcaaaa taaaggtatt tgtataaaaa aaaaaaaaaa 1320
aaactcga                                                         1328
```

<210> 25
<211> 1758
<212> DNA
<213> Homo sapiens

<400> 25
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gttttttttt tttttttttt aaagagttgc aacaattcat ctttattttct tattttcctc 60
tggagatgca gaatttggta tatttcaccc caagtatatt tgggatagtt ggctcctcgc 120
tgggtcagga tggctgggtg ccttctcccc tggcatggtt ctcttctctg cagggcgagg 180
ggcagggagc tagtaaaacc tcgcaatgac agccgcaatg gcagacccaa tggagcccag 240
gatgaacttg gtcaatccgg agagtccagt tgctcccagt gactgcagag tagccacaag 300
gctgcccgag gcaactccac ccccattggc aatggccgcc gcggacatca tcttggctgc 360
tatggaggac gaggccgattc ccgccgcagt gaagcccatg gcactgagtg gcggcggtgg 420
atatccgaga caatctgctg ggaatttctt gggttgacag ctcttggatc cctattttga 480
acagtggtag tgtcctggat tactttttcag aaagaagtaa tcctttttat gacagaacat 540
gtaataatga agtggtcaaa atgcagaggc taacattaga acacttgaat cagatggttg 600
gaatcgagta catccttttg catgctcaag agcccattct tttcatcatt cggaagcaac 660
agcggcagtc ccctgcccaa gttatcccac tagctgatta ctatatcatt gctggagtga 720
tctatcaggc accagacttg ggatcagtta taaactctag agtgcttact gcagtgcatg 780
gtattcagtc agcttttgat gaagctatgt catactgtcg atatcatcct tccaaagggt 840
attggtggca cttcaaagat catgaagagc aagataaagt cagacctaaa gccaaaagga 900
aagaagaacc aagctctatt tttcagagac aacgtgtgga tgctttactt ttagacctca 960
gacaaaaatt tccacccaaa tttgtgcagc taaagcctgg agaaaagcct gttccagtgg 1020
atcaaacaaa gaaagaggca gaacctatac cagaaactgt aaaacctgag gagaaggaga 1080
ccacaaagaa tgtacaacag acagtgagtg ctaaaggccc ccctgaaaaa cggatgagac 1140
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cagtactgtg gctcttgagc tttgaagtac tttattgtaa ccttcttatt tgtatggaat 1260
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gcccatagtc ccagctactc gggaggctga cacgagagac ttgcttgaac ccgggaggtg 1500
gaggttgccc tgagctgatt atcatgctgt tgcactccag cttgggcgac agagcgagac 1560
tttgtctcaa aaagaagaa aagatattat tcccatcatg atttcttgtg aatatttgtt 1620
atatgtcttc tgttaccttt cctctcccgg aattgagcaa cctacacact cacatgttta 1680
ctggtagata tgtttaaaag caaataaagg tattggtata tattgcttca aaaaaaaaaa 1740
aaaaaaaaaa aactcgag                                              1758
```

<210> 26
<211> 493
<212> DNA
<213> Homo sapiens

<400> 26
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gaggcgagcg gcagggcctg gtggcgagag cgcggctgtc actgcgcccg agcatcccag 60
agctttccga gcggacgagc cggccgtgcc gggcatcccc agcctcgcta ccctcgcagc 120
```

```
acacgtcgag ccccgcacag gcaagggtcc ggaacttagc ccaaagcacg tttcccctgg 180
cagcgcagga gacgcccggc cgcgcgccgg cgcacgcccc cctctcctcc tttgttccgg 240
gggtcggcgg ccgctctcct gccagcgtcg ggatctcggc cccgggaggc gggccgtcgg 300
gcgcagccgc gaagattccg ttggaactga cgcagagccg agtgcagaag atctgggtgc 360
ccgtggacca caggccctcg ttgcccagat cctgtgggcc aaagctgacc aactcccccg 420
ccgtcttcgt catggtgggc ctcccccgcc cggggcaaga cctacttctc cacgaaagct 480
tactcgctgc ctc                                                    493
```

```
<210> 27
<211> 1331
<212> DNA
<213> Homo sapiens
```

```
<400> 27
ggtggatatc cgagacaatc tgctgggaat ttcttgggtt gacagctctt ggatccctat 60
tttgaacagt ggtagtgtcc tggattactt ttcagaaaga agtaatcctt tttatgacag 120
aacatgtaat aatgaagtgg tcaaaatgca gaggctaaca ttagaacact tgaatcagat 180
ggttggaatc gagtacatcc ttttgcatgc tcaagagccc attcttttca tcattcggaa 240
gcaacagcgg cagtcccctg cccaagttat cccactagct gattactata tcattgctgg 300
agtgatctat caggcaccag acttgggatc agttataaac tctagagtgc ttactgcagt 360
gcatggtatt cagtcagctt ttgatgaagc tatgtcatac tgtcgatatc atccttccaa 420
agggtattgg tggcacttca aagatcatga agagcaagat aaagtcagac ctaaagccaa 480
aaggaaagaa gaaccaagct ctattttca gagacaacgt gtggatgctt tactttttaga 540
cctcagacaa aaatttccac ccaaatttgt gcagctaaag cctggagaaa agcctgttcc 600
agtggatcaa acaaagaaag aggcagaacc tataccagaa actgtaaaac ctgaggagaa 660
ggagaccaca aagaatgtac aacagacagt gagtgctaaa ggcccccctg aaaaacggat 720
gagacttcag tgagtactgg acaaaagaga agcctggaag actcctcatg ctagttatca 780
tacctcagta ctgtggctct tgagctttga agtactttat tgtaaccttc ttatttgtat 840
ggaatgcgct tatttttga aaggatatta ggccggatgt ggtggctcac gcctgtaatc 900
ccagcacttt gggaggccat ggcgggtgga tcacttgagg tcagaagttc aagaccagcc 960
tgaccaatat ggtgaaaccc cgtctctact aaaaatacaa aaattagccg ggcgtggtgg 1020
cgggcgccca tagtcccagc tactcgggag gctgagacag gagacttgct tgaacccggg 1080
aggtggaggt tgccctgagc tgattatcat gctgttgcac tccagcttgg cgacagagc 1140
gagactttgt ctcaaaaaaa gaagaaaaga tattattccc atcatgattt cttgtgaata 1200
tttgttatat gtcttctgta acctttcctc tcccggactt gagcaaccta cacactcaca 1260
tgtttactgg tagatatgtt taaaagcaaa ataaaggtat tggtataaaa aaaaaaaaaa 1320
aaaaactcga g                                                     1331
```

```
<210> 28
<211> 1333
<212> DNA
<213> Homo sapiens
```

```
<400> 28
cggcggtgga tatccgagac aatctgctgg gaatttcttg ggttgacagc tcttggatcc 60
ctattttgaa cagtggtagt gtcctggatt acttttcaga agaagtaat cctttttatg 120
acagaacatg taataatgaa gtggtcaaaa tgcagaggct aacattagaa cacttgaatc 180
agatggttgg aatcgagtac atccttttgc atgctcaaga gcccattctt ttcatcattc 240
ggaagcaaca gcggcagtcc cctgcccaag ttatcccact agctgattac tatatcattg 300
ctggagtgat ctatcaggca ccagacttgg gatcagttat aaactctaga gtgcttactg 360
cagtgcatgg tatt cagtca gcttttgatg aagctatgtc atactgtcga tatcatcctt 420
ccaaagggta ttggtggcac ttcaaagatc atgaagagca agataaagtc agacctaaag 480
ccaaaaggaa agaagaacca agctctattt tcagagaca cgtgtggat gctttacttt 540
tagacctcag acaaaaattt ccacccaaat ttgtgcagct aaagcctgga gaaaagcctg 600
ttccagtgga tcaaacaaag aaagaggcag aacctatacc agaaactgta aaacctgagg 660
```

```
agaaggagac cacaaagaat gtacaacaga cagtgagtgc taaaggcccc cctgaaaaac 720
ggatgagact tcagtgagta ctggacaaaa gagaagcctg gaagactcct catgctagtt 780
atcatacctc agtactgtgg ctcttgagct ttgaagtact ttattgtaac cttcttattt 840
gtatggaatg cgcttatttt ttgaaaggat attaggccgg atgtggtggc tcacgcctgt 900
aatcccagca ctttgggagg ccatggcggg tggatcactt gaggtcagaa gttcaagacc 960
agcctgacca atatggtgaa acccgtctc tactaaaaat acaaaaatta gccgggcgtg 1020
gtggcgggcg cccatagtcc cagctactcg ggaggctgag acaggagact tgcttgaacc 1080
cgggaggtgg aggttgccct gagctgatta tcatgctgtt gcactccagc ttgggcgaca 1140
gagcgagact ttgtctcaaa aaagaagaaa agatattatt cccatcatga tttcttgtga 1200
atatttgtga tatgtcttct gtaacctttc ctctcccgga cttgagcaac ctacacactc 1260
acatgtttac tggtagatat gtttaaaagc aaaataaagg tatttgtata aaaaaaaaaa 1320
aaaaaaactc gag                                                     1333
```

<210> 29
<211> 813
<212> DNA
<213> Homo sapiens

<400> 29
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ctgagctgca cttcagcgaa ttcacctcgg ctgtggctga catgaagaac tccgtggcgg 60
accgagacaa cagccccagc tcctgtgctg gcctcttcat tgcttcacac atcgggtttg 120
actggcccgg ggtctgggtc cacctggaca tcgctgctcc agtgcatgct ggcgagcgag 180
ccacaggctt tggggtggct ctcctactgg ctcttttttgg ccgtgcctcc gaggacccgc 240
tgctgaacct ggtatccccg ctggactgtg aggtggatgc ccaggaaggc gacaacatgg 300
ggcgtgactc caagagacgg aggctcgtgt gagggctact tcccagctgg tgacacaggg 360
ttccttacct cattttgcac tgactgattt taagcaattg aaagattaac taactcttaa 420
gatgagtttg gcttctcctt ctgtgcccag tggtgacagg agtgagccat tcttctctta 480
gaagcagctt aggggcttgg tggggtctgg agaaaattgt cacagacccc ataggtctcc 540
atctgtaagc tctgtccctt gtcctccacc ctggtctttta gagccacctc aggtcaccct 600
ctgtagtgag tgtacttcct gacccaggcc cttgctcaag ctggggctcc ctggggtgtc 660
taaccagccc tgggtagatg tgactggctg ttaggaccc cattctgtga agcaggagac 720
cctcacagct cccaccaacc cccagttcac ttgaagttga attaaatatg gccacaacat 780
aaaaaaaaaa aaaaaaaaaa aaaaaactc gag                                813
```

<210> 30
<211> 1316
<212> DNA
<213> Homo sapiens

<400> 30
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caggcgccca gtcatggccc aagagacagc accaccgtgt ggcccagtct caaggggtga 60
cagtccaatc atagaaaaga tggaaaaaag gacatgtgcc ctgtgccctg aaggccacga 120
gtggagtcaa atatactttt caccatcagg aaatatagtt gctcatgaaa actgtttgct 180
gtattcatca ggactggtgg agtgtgagac tcttgatcta cgtaatacaa ttagaaactt 240
tgatgtcaaa tctgtaaaga aagagatctg gagaggaaga agattgaaat gctcattctg 300
taacaaagga ggcgccaccg tggggtgtga tttatggttc tgtaagaaga gttaccacta 360
tgtctgtgcc aaaaaggacc aagcaattct tcaagttgat ggaaaccatg gaacttacaa 420
attattttgc ccagaacatt ctccagaaca agaagaggcc actgaaagtg ctgatgaccc 480
aagcatgaag aagaagagag gaaaaaacaa acgcctctca tcaggccctc ctgcacagcc 540
aaaaacgatg aaatgtagta acgccaaaag acatatgaca gaagagcctc atggtcacac 600
agatgcagct gtcaaatctc cttttcttaa gaaatgccag gaagcaggac ttcttactga 660
actatttgaa cacatactag aaaatatgga ttcagttcat ggaagacttg tggatgagac 720
tgcctcagag tcggactatg aagggatcga gaccttactg tttgactgtg gattattaa 780
agacacacta agaaaattcc aagaagtaat caagagtaaa gcttgtgaat gggaagaaag 840
gcaaaggcag atgaagcagc agcttgaggc acttgcagac ttacaacaaa gcttgtgctc 900
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12

```
atttcaagaa aatggggacc tggactgctc aagttctaca tcaggatcct tgctacctcc 960
tgaggaccac cagtggaaagc tgttcctcag gaaaactgga tggggcctcc atgttctcca 1020
aggatcgagg aagtcttcct gcctaccctg cccaccccag tcaagggcag caacaccaga 1080
gctttgctca gccttaaatg gaatcttaga gctttctctt gcttctgcta ctcctacaga 1140
tggcctcatc atggtctcca ctcagtatta ataactccat cagcatagag caaactcaac 1200
actgtgcatt gcacactgtt accatgggtt tatgctcact atcatatcac attgccaata 1260
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```

<210> 31
<211> 1355
<212> DNA
<213> Homo sapiens

<400> 31
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cggcggtgga tatccgagac aatctgctgg gaatttcttg ggttgacagc tcttggatcc 60
ctattttgaa cagtggtagt gtcctggatt acttttcaga aagaagtaat cctttttatg 120
acagaacatg taataatgaa gtggtcaaaa tgcagaggct aacattagaa cacttgaatc 180
agatggttgg aatcgagtac atccttttgc atgctcaaga gcccattctt ttcatcattc 240
ggaagcaaca gcggcagtcc cctgcccaag ttatcccact agctgattac tatatcattg 300
ctggagtgat ctatcaggca ccagacttgg gatcagttat aaactctaga gtgcttactg 360
cagtgcatgg tattcagtca gcttttgatg aagctatgtc atactgtcga tatcatcctt 420
ccaaagggta ttggtggcac ttcaaagatc atgaagagca agataaagtc agacctaaag 480
ccaaaaggaa agaagaacca agctctattt ttcagagaca acgtgtggat gctttactt 540
tagacctcag acaaaaattt ccacccaaat ttgtgcagct aaagcctgga gaaaagcctg 600
ttccagtgga tcaaacaaag aaagaggcag aacctatacc agaaactgta aaacctgagg 660
agaaggagac cacaaagaat gtacaacaga cagtgagtgc taaaggcccc cctgaaaaac 720
ggatgagact tcagtgagta ctggacaaaa gagaagcctg gaagactcct catgctagtt 780
atcataacctc agtactgtgg ctcttgagct ttgaagtact ttattgtaac cttcttattt 840
gtatggaatg cgcttatttt ttgaaaggat attaggccgg atgtggtggc tcacgcctgt 900
aatcccagca ctttgggagg ccatggcggg tggatcactt gaggtcagaa gttcaagacc 960
agcctgacca atatggtgaa accccgtctc tactaaaaat acaaaaatta gccgggcgtg 1020
gtggcgggcg cccatagtcc cagctactcg ggaggctgag acaggagact tgcttgaacc 1080
cgggaggtgg aggttgccct gagctgatta tcatgctgtt gcactccagc ttgggcgaca 1140
gaacgagact ttgtctcaaa aaagaagaa aagatattat tcccatcatg atttcttgtg 1200
aatatttgtt atatgtcttc tggtaacctt tcctctcccg gacttgaagc aacctcacac 1260
actcacatgt ttactggtag atatgtttta aaagcaaaat aaaggtattt gtttttccaa 1320
aaaaaaaaaa aaaaaaaaaa aaaaaaaaac tcgag                           1355
```

<210> 32
<211> 80
<212> PRT
<213> Homo sapiens

<400> 32
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Val Ser Arg Ile Arg Gly Gly Ala Lys Lys Arg Lys Lys Lys Ser Tyr
 1               5                  10                  15

Thr Thr Pro Lys Lys Asp Lys His Gln Arg Lys Lys Val Gln Pro Ala
            20                  25                  30

Val Leu Lys Tyr Tyr Lys Val Asp Glu Asn Gly Lys Ile Ser Cys Leu
        35                  40                  45

Arg Arg Glu Cys Pro Ser Asp Glu Cys Gly Ala Gly Val Phe Met Ala
    50                  55                  60
```

Ser His Phe Asp Arg His Tyr Cys Gly Lys Cys Cys Leu Thr His Cys
65   70   75   80

<210> 33
<211> 130
<212> PRT
<213> Homo sapiens

<400> 33
Glu Ile Ser Asn Glu Val Arg Lys Phe Arg Thr Leu Thr Glu Leu Ile
1   5   10   15

Leu Asp Ala Gln Glu His Val Lys Asn Pro Tyr Lys Gly Lys Lys Leu
            20   25   30

Lys Lys His Pro Asp Phe Pro Lys Lys Pro Leu Thr Pro Tyr Phe Arg
        35   40   45

Phe Phe Met Glu Lys Arg Ala Lys Tyr Ala Lys Leu His Pro Gln Met
    50   55   60

Ser Asn Leu Asp Leu Thr Lys Ile Leu Ser Lys Lys Tyr Lys Glu Leu
65   70   75   80

Pro Glu Lys Lys Lys Met Lys Tyr Val Pro Asp Phe Gln Arg Arg Glu
            85   90   95

Thr Gly Val Arg Ala Lys Pro Gly Pro Ile Gln Gly Gly Ser Pro Pro
            100   105   110

Pro Tyr Pro Glu Cys Gln Glu Ser Asp Ile Pro Glu Lys Pro Gln Asp
        115   120   125

Pro Pro
    130

<210> 34
<211> 506
<212> PRT
<213> Homo sapiens

<400> 34
Asn Ser Glu Lys Glu Ile Pro Val Leu Asn Glu Leu Pro Val Pro Met
1   5   10   15

Val Ala Arg Tyr Ile Arg Ile Asn Pro Gln Ser Trp Phe Asp Asn Gly
            20   25   30

Ser Ile Cys Met Arg Met Glu Ile Leu Gly Cys Pro Leu Pro Asp Pro

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```
        35                     40                      45
Asn Asn Tyr Tyr His Arg Arg Asn Glu Met Thr Thr Thr Asp Asp Leu
    50                  55                  60

Asp Phe Lys His His Asn Tyr Lys Glu Met Arg Gln Leu Met Lys Val
   65                  70                  75                  80

Val Asn Glu Met Cys Pro Asn Ile Thr Arg Ile Tyr Asn Ile Gly Lys
                85                  90                      95

Ser His Gln Gly Leu Lys Leu Tyr Ala Val Glu Ile Ser Asp His Pro
                100                 105                 110

Gly Glu His Glu Val Gly Glu Pro Glu Phe His Tyr Ile Ala Gly Ala
            115                 120                 125

His Gly Asn Glu Val Leu Gly Arg Glu Leu Leu Leu Leu Leu Leu His
    130                 135                 140

Phe Leu Cys Gln Glu Tyr Ser Ala Gln Asn Ala Arg Ile Val Arg Leu
145                 150                 155                 160

Val Glu Glu Thr Arg Ile His Ile Leu Pro Ser Leu Asn Pro Asp Gly
                165                 170                 175

Tyr Glu Lys Ala Tyr Glu Gly Gly Ser Glu Leu Gly Gly Trp Ser Leu
            180                 185                 190

Gly Arg Trp Thr His Asp Gly Ile Asp Ile Asn Asn Asn Phe Pro Asp
    195                 200                 205

Leu Asn Ser Leu Leu Trp Glu Ala Glu Asp Gln Gln Asn Ala Pro Arg
    210                 215                 220

Lys Val Pro Asn His Tyr Ile Ala Ile Pro Glu Trp Phe Leu Ser Glu
225                 230                 235                 240

Asn Ala Thr Val Ala Thr Glu Thr Arg Ala Val Ile Ala Trp Met Glu
                245                 250                 255

Lys Ile Pro Phe Val Leu Gly Gly Asn Leu Gln Gly Gly Glu Leu Val
                260                 265                 270

Val Ala Tyr Pro Tyr Asp Met Val Arg Ser Leu Trp Lys Thr Gln Glu
            275                 280                 285

His Thr Pro Thr Pro Asp Asp His Val Phe Arg Trp Leu Ala Tyr Ser
    290                 295                 300

Tyr Ala Ser Thr His Arg Leu Met Thr Asp Ala Arg Arg Arg Val Cys
305                 310                 315                 320

His Thr Glu Asp Phe Gln Lys Glu Glu Gly Thr Val Asn Gly Ala Ser
            325                 330                 335
```

Trp His Thr Val Ala Gly Ser Leu Asn Asp Phe Ser Tyr Leu His Thr
       340   345   350

Asn Cys Phe Glu Leu Ser Ile Tyr Val Gly Cys Asp Lys Tyr Pro His
       355   360   365

Glu Ser Glu Leu Pro Glu Glu Trp Glu Asn Asn Arg Glu Ser Leu Ile
    370   375   380

Val Phe Met Glu Gln Val His Arg Gly Ile Lys Gly Ile Val Arg Asp
385   390   395   400

Leu Gln Gly Lys Gly Ile Ser Asn Ala Val Ile Ser Val Glu Gly Val
         405   410   415

Asn His Asp Ile Arg Thr Ala Ser Asp Gly Asp Tyr Trp Arg Leu Leu
       420   425   430

Asn Pro Gly Glu Tyr Val Val Thr Ala Lys Ala Glu Gly Phe Ile Thr
       435   440   445

Ser Thr Lys Asn Cys Met Val Gly Tyr Asp Met Gly Ala Thr Arg Cys
      450   455   460

Asp Phe Thr Leu Thr Lys Thr Asn Leu Ala Arg Ile Arg Glu Ile Met
465   470   475   480

Glu Thr Phe Gly Lys Gln Pro Val Ser Leu Pro Ser Arg Arg Leu Lys
         485   490   495

Leu Arg Gly Arg Lys Arg Arg Gln Arg Gly
       500   505

<210> 35
<211> 96
<212> PRT
<213> Homo sapiens

<400> 35
Met Asn Gly Glu Ala Asp Cys Pro Thr Asp Leu Glu Met Ala Ala Pro
  1   5   10   15

Arg Gly Gln Asp Arg Trp Ser Gln Glu Asp Met Leu Thr Leu Leu Glu
      20   25   30

Cys Met Lys Asn Asn Leu Pro Ser Asn Asp Ser Ser Gln Phe Lys Thr
      35   40   45

Thr Gln Thr His Met Asp Arg Glu Lys Val Ala Leu Lys Asp Phe Ser
    50   55   60

Gly Asp Met Cys Lys Leu Lys Trp Val Glu Ile Ser Asn Glu Val Arg
65   70   75   80

Lys Phe Arg Thr Leu Thr Glu Leu Ile Leu Asp Thr Gln Glu His Val
                85   90   95

<210> 36
<211> 129
<212> PRT
<213> Homo sapiens

<400> 36
Gly Ile Val Val Phe Ser Leu Gly Ser Met Val Ser Glu Ile Pro Glu
1   5   10   15

Lys Lys Ala Val Ala Ile Ala Asp Ala Leu Gly Lys Ile Pro Gln Thr
            20   25   30

Val Leu Trp Arg Tyr Thr Gly Thr Arg Pro Ser Asn Leu Ala Asn Asn
        35   40   45

Thr Ile Leu Val Gln Trp Leu Pro Gln Asn Asp Leu Leu Gly His Pro
    50   55   60

Met Thr Arg Ala Phe Ile Thr His Ala Ser Ser His Gly Val Asn Glu
65   70   75   80

Ser Ile Cys Asn Gly Val Pro Met Val Met Ile Pro Leu Phe Gly Asp
                85   90   95

Gln Met Asp Asn Ala Lys Arg Arg Glu Thr Lys Gly Ala Gly Val Thr
            100   105   110

Leu Asn Val Leu Glu Met Thr Ser Glu Asp Leu Glu Asp Ala Leu Lys
        115   120   125

Ser

<210> 37
<211> 238
<212> PRT
<213> Homo sapiens

<400> 37
Asn Leu Leu Gly Ile Ser Trp Val Asp Ser Ser Trp Ile Pro Ile Leu
1   5   10   15

Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser Glu Arg Ser Asn Pro Phe
            20   25   30

Tyr Asp Arg Thr Cys Asn Asn Glu Val Val Lys Met Gln Arg Leu Thr
        35   40   45

Leu Glu His Leu Asn Gln Met Val Gly Ile Glu Tyr Ile Leu Leu His
        50   55   60

```
Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg Lys Gln Gln Arg Gln Ser
65                  70                  75                  80

Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr Tyr Ile Ile Ala Gly Val
                 85                  90                  95

Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val Ile Asn Ser Arg Val Leu
             100                 105                 110

Thr Ala Val His Gly Ile Gln Ser Ala Phe Asp Glu Ala Met Ser Tyr
             115                 120                 125

Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp Trp His Phe Lys Asp His
         130                 135                 140

Glu Glu Gln Asp Lys Val Arg Pro Lys Ala Lys Arg Lys Glu Glu Pro
145                 150                 155                 160

Ser Ser Ile Phe Gln Arg Gln Arg Val Asp Ala Leu Leu Leu Asp Leu
             165                 170                 175

Arg Gln Lys Phe Pro Pro Lys Phe Val Gln Leu Lys Pro Gly Glu Lys
             180                 185                 190

Pro Val Pro Val Asp Gln Thr Lys Lys Glu Ala Glu Pro Ile Pro Glu
         195                 200                 205

Thr Val Lys Pro Glu Glu Lys Glu Thr Thr Lys Asn Val Gln Gln Thr
         210                 215                 220

Val Ser Ala Lys Gly Pro Pro Glu Lys Arg Met Arg Leu Gln
225                 230                 235
```

<210> 38
<211> 202
<212> PRT
<213> Homo sapiens

<400> 38
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Lys Gly Ser Glu Gly Glu Asn Pro Leu Thr Val Pro Gly Arg Glu Lys
  1               5                  10                  15

Glu Gly Met Leu Met Gly Val Lys Pro Gly Glu Asp Ala Ser Gly Pro
             20                  25                  30

Ala Glu Asp Leu Val Arg Arg Ser Glu Lys Asp Thr Ala Ala Val Val
         35                  40                  45

Ser Arg Gln Gly Ser Ser Leu Asn Leu Phe Glu Asp Val Gln Ile Thr
     50                  55                  60

Glu Pro Glu Ala Glu Pro Glu Ser Lys Ser Glu Pro Arg Pro Pro Ile
65                  70                  75                  80
```

Ser Ser Pro Arg Ala Pro Gln Thr Arg Ala Val Lys Pro Arg Leu His
            85   90   95

Pro Val Lys Pro Met Asn Ala Thr Ala Thr Lys Val Ala Asn Cys Ser
            100   105   110

Leu Gly Thr Ala Thr Ile Ile Gly Glu Asn Leu Asn Asn Glu Val Met
            115   120   125

Met Lys Lys Tyr Ser Pro Ser Asp Pro Ala Phe Ala Tyr Ala Gln Leu
    130   135   140

Thr His Asp Glu Leu Ile Gln Leu Val Leu Lys Gln Lys Glu Thr Ile
145   150   155   160

Ser Lys Lys Glu Phe Gln Val Arg Glu Leu Glu Asp Tyr Ile Asp Asn
            165   170   175

Leu Leu Val Arg Val Met Glu Glu Thr Pro Asn Ile Leu Arg Ile Pro
            180   185   190

Thr Gln Val Gly Lys Lys Ala Gly Lys Met
            195   200

<210> 39
<211> 243
<212> PRT
<213> Homo sapiens

<400> 39
Val Asn Ala Leu Gly Ile Met Ala Ala Val Asp Ile Arg Asp Asn Leu
    1   5   10   15

Leu Gly Ile Ser Trp Val Asp Ser Ser Trp Ile Pro Ile Leu Asn Ser
            20   25   30

Gly Ser Val Leu Asp Tyr Phe Ser Glu Arg Ser Asn Pro Phe Tyr Asp
            35   40   45

Arg Thr Cys Asn Asn Glu Val Val Lys Met Gln Arg Leu Thr Leu Glu
    50   55   60

His Leu Asn Gln Met Val Gly Ile Glu Tyr Ile Leu Leu His Ala Gln
65   70   75   80

Glu Pro Ile Leu Phe Ile Ile Arg Lys Gln Gln Arg Gln Ser Pro Ala
            85   90   95

Gln Val Ile Pro Leu Ala Asp Tyr Tyr Ile Ile Ala Gly Val Ile Tyr
            100   105   110

Gln Ala Pro Asp Leu Gly Ser Val Ile Asn Ser Arg Val Leu Thr Ala
            115   120   125

```
Val His Gly Ile Gln Ser Ala Phe Asp Glu Ala Met Ser Tyr Cys Arg
    130                 135             140

Tyr His Pro Ser Lys Gly Tyr Trp Trp His Phe Lys Asp His Glu Glu
145                 150             155             160

Gln Asp Lys Val Arg Pro Lys Ala Lys Arg Lys Glu Glu Pro Ser Ser
                165             170             175

Ile Phe Gln Arg Gln Arg Val Asp Ala Leu Leu Leu Asp Leu Arg Gln
            180             185             190

Lys Ile Ser Thr Gln Ile Cys Ala Val Asp Gln Thr Lys Lys Glu Ala
            195             200             205

Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr Thr Lys
    210             215             220

Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys Arg Met
225             230             235             240

Arg Leu Gln
```

<210> 40
<211> 245
<212> PRT
<213> Homo sapiens

<400> 40
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Ala Ala Val Asp Ile Arg Asp Asn Leu Leu Gly Ile Ser Trp Val Asp
  1           5               10                  15

Ser Ser Trp Ile Pro Ile Leu Asn Ser Gly Ser Val Leu Asp Tyr Phe
            20              25                  30

Ser Glu Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val
        35              40                  45

Val Lys Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly
    50                  55                  60

Ile Glu Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile
65                  70                  75                  80

Arg Lys Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp
                85              90                  95

Tyr Tyr Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser
            100             105             110

Val Ile Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala
        115             120             125
```

```
Phe Asp Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr
    130                 135                 140

Trp Trp His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys
145                 150                 155                 160

Ala Lys Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val
                165                 170                 175

Asp Ala Leu Leu Leu Asp Leu Arg Gln Lys Phe Pro Pro Lys Phe Val
                180                 185                 190

Gln Leu Lys Pro Gly Glu Lys Pro Val Pro Val Asp Gln Thr Lys Lys
            195                 200                 205

Glu Ala Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr
    210                 215                 220

Thr Lys Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys
225                 230                 235                 240

Arg Met Arg Leu Gln
                245


<210> 41
<211> 163
<212> PRT
<213> Homo sapiens

<400> 41
Gly Glu Arg Gln Gly Leu Val Ala Arg Ala Arg Leu Ser Leu Arg Pro
    1               5               10                  15

Ser Ile Pro Glu Leu Ser Glu Arg Thr Ser Arg Pro Cys Arg Ala Ser
            20                  25                  30

Pro Ala Ser Leu Pro Ser Gln His Thr Ser Ser Pro Ala Gln Ala Arg
            35                  40                  45

Val Arg Asn Leu Ala Gln Ser Thr Phe Pro Leu Ala Ala Gln Glu Thr
    50                  55                  60

Pro Gly Arg Ala Pro Ala His Ala Pro Leu Ser Ser Phe Val Pro Gly
65                  70                  75                  80

Val Gly Gly Arg Ser Pro Ala Ser Val Gly Ile Ser Ala Pro Gly Gly
                85                  90                  95

Gly Pro Ser Gly Ala Ala Ala Lys Ile Pro Leu Glu Leu Thr Gln Ser
            100                 105                 110

Arg Val Gln Lys Ile Trp Val Pro Val Asp His Arg Pro Ser Leu Pro
    115                 120                 125

Arg Ser Cys Gly Pro Lys Leu Thr Asn Ser Pro Ala Val Phe Val Met
```

                 130   135   140

    Val Gly Leu Pro Arg Pro Gly Gln Asp Leu Leu Leu His Glu Ser Leu
    145   150   155   160

    Leu Ala Ala

    <210> 42
    <211> 243
    <212> PRT
    <213> Homo sapiens

    <400> 42
    Val Asp Ile Arg Asp Asn Leu Leu Gly Ile Ser Trp Val Asp Ser Ser
     1   5   10   15

    Trp Ile Pro Ile Leu Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser Glu
                  20   25   30

    Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val Val Lys
            35   40   45

    Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly Ile Glu
          50   55   60

    Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg Lys
    65   70   75   80

    Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr Tyr
                  85   90   95

    Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val Ile
                  100   105   110

    Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala Phe Asp
            115   120   125

    Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp Trp
          130   135   140

    His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys Ala Lys
    145   150   155   160

    Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val Asp Ala
                  165   170   175

    Leu Leu Leu Asp Leu Arg Gln Lys Phe Pro Pro Lys Phe Val Gln Leu
                  180   185   190

    Lys Pro Gly Glu Lys Pro Val Pro Val Asp Gln Thr Lys Lys Glu Ala
            195   200   205

    Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr Thr Lys
          210   215   220

Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys Arg Met
225   230   235   240

Arg Leu Gln

<210> 43
<211> 244
<212> PRT
<213> Homo sapiens

<400> 43
Ala Val Asp Ile Arg Asp Asn Leu Leu Gly Ile Ser Trp Val Asp Ser
  1   5   10   15

Ser Trp Ile Pro Ile Leu Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser
          20   25   30

Glu Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val Val
          35   40   45

Lys Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly Ile
      50   55   60

Glu Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg
65   70   75   80

Lys Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr
              85   90   95

Tyr Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val
          100   105   110

Ile Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala Phe
          115   120   125

Asp Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp
      130   135   140

Trp His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys Ala
145   150   155   160

Lys Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val Asp
              165   170   175

Ala Leu Leu Leu Asp Leu Arg Gln Lys Phe Pro Pro Lys Phe Val Gln
              180   185   190

Leu Lys Pro Gly Glu Lys Pro Val Pro Val Asp Gln Thr Lys Lys Glu
          195   200   205

Ala Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr Thr
      210   215   220

```
Lys Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys Arg
225             230             235             240

Met Arg Leu Gln


<210> 44
<211> 109
<212> PRT
<213> Homo sapiens

<400> 44
Glu Leu His Phe Ser Glu Phe Thr Ser Ala Val Ala Asp Met Lys Asn
  1             5               10              15

Ser Val Ala Asp Arg Asp Asn Ser Pro Ser Ser Cys Ala Gly Leu Phe
            20              25              30

Ile Ala Ser His Ile Gly Phe Asp Trp Pro Gly Val Trp Val His Leu
         35              40              45

Asp Ile Ala Ala Pro Val His Ala Gly Glu Arg Ala Thr Gly Phe Gly
     50              55              60

Val Ala Leu Leu Leu Ala Leu Phe Gly Arg Ala Ser Glu Asp Pro Leu
  65              70              75              80

Leu Asn Leu Val Ser Pro Leu Asp Cys Glu Val Asp Ala Gln Glu Gly
            85              90              95

Asp Asn Met Gly Arg Asp Ser Lys Arg Arg Arg Leu Val
            100             105


<210> 45
<211> 324
<212> PRT
<213> Homo sapiens

<400> 45
Arg Arg Pro Val Met Ala Gln Glu Thr Ala Pro Pro Cys Gly Pro Val
  1             5               10              15

Ser Arg Gly Asp Ser Pro Ile Ile Glu Lys Met Glu Lys Arg Thr Cys
            20              25              30

Ala Leu Cys Pro Glu Gly His Glu Trp Ser Gln Ile Tyr Phe Ser Pro
         35              40              45

Ser Gly Asn Ile Val Ala His Glu Asn Cys Leu Leu Tyr Ser Ser Gly
     50              55              60

Leu Val Glu Cys Glu Thr Leu Asp Leu Arg Asn Thr Ile Arg Asn Phe
  65              70              75              80
```

```
Asp Val Lys Ser Val Lys Lys Glu Ile Trp Arg Gly Arg Arg Leu Lys
            85                    90                    95

Cys Ser Phe Cys Asn Lys Gly Gly Ala Thr Val Gly Cys Asp Leu Trp
            100               105               110

Phe Cys Lys Lys Ser Tyr His Tyr Val Cys Ala Lys Lys Asp Gln Ala
        115               120               125

Ile Leu Gln Val Asp Gly Asn His Gly Thr Tyr Lys Leu Phe Cys Pro
    130                   135               140

Glu His Ser Pro Glu Gln Glu Glu Ala Thr Glu Ser Ala Asp Asp Pro
145               150               155               160

Ser Met Lys Lys Lys Arg Gly Lys Asn Lys Arg Leu Ser Ser Gly Pro
            165               170               175

Pro Ala Gln Pro Lys Thr Met Lys Cys Ser Asn Ala Lys Arg His Met
            180               185               190

Thr Glu Glu Pro His Gly His Thr Asp Ala Ala Val Lys Ser Pro Phe
        195               200               205

Leu Lys Lys Cys Gln Glu Ala Gly Leu Leu Thr Glu Leu Phe Glu His
    210               215               220

Ile Leu Glu Asn Met Asp Ser Val His Gly Arg Leu Val Asp Glu Thr
225               230               235               240

Ala Ser Glu Ser Asp Tyr Glu Gly Ile Glu Thr Leu Leu Phe Asp Cys
            245               250               255

Gly Leu Phe Lys Asp Thr Leu Arg Lys Phe Gln Glu Val Ile Lys Ser
        260               265               270

Lys Ala Cys Glu Trp Glu Glu Arg Gln Arg Gln Met Lys Gln Gln Leu
        275               280               285

Glu Ala Leu Ala Asp Leu Gln Gln Ser Leu Cys Ser Phe Gln Glu Asn
    290               295               300

Gly Asp Leu Asp Cys Ser Ser Ser Thr Ser Gly Ser Leu Leu Pro Pro
305               310               315               320

Glu Asp His Gln
```

```
<210> 46
<211> 244
<212> PRT
<213> Homo sapiens

<400> 46
Ala Val Asp Ile Arg Asp Asn Leu Leu Gly Ile Ser Trp Val Asp Ser
```

```
      1               5              10              15

  Ser Trp Ile Pro Ile Leu Asn Ser Gly Ser Val Leu Asp Tyr Phe Ser
              20              25              30

  Glu Arg Ser Asn Pro Phe Tyr Asp Arg Thr Cys Asn Asn Glu Val Val
          35              40              45

  Lys Met Gln Arg Leu Thr Leu Glu His Leu Asn Gln Met Val Gly Ile
      50              55              60

  Glu Tyr Ile Leu Leu His Ala Gln Glu Pro Ile Leu Phe Ile Ile Arg
  65              70              75              80

  Lys Gln Gln Arg Gln Ser Pro Ala Gln Val Ile Pro Leu Ala Asp Tyr
              85              90              95

  Tyr Ile Ile Ala Gly Val Ile Tyr Gln Ala Pro Asp Leu Gly Ser Val
              100             105             110

  Ile Asn Ser Arg Val Leu Thr Ala Val His Gly Ile Gln Ser Ala Phe
          115             120             125

  Asp Glu Ala Met Ser Tyr Cys Arg Tyr His Pro Ser Lys Gly Tyr Trp
      130             135             140

  Trp His Phe Lys Asp His Glu Glu Gln Asp Lys Val Arg Pro Lys Ala
  145             150             155             160

  Lys Arg Lys Glu Glu Pro Ser Ser Ile Phe Gln Arg Gln Arg Val Asp
              165             170             175

  Ala Leu Leu Leu Asp Leu Arg Gln Lys Phe Pro Pro Lys Phe Val Gln
              180             185             190

  Leu Lys Pro Gly Glu Lys Pro Val Pro Val Asp Gln Thr Lys Lys Glu
          195             200             205

  Ala Glu Pro Ile Pro Glu Thr Val Lys Pro Glu Glu Lys Glu Thr Thr
      210             215             220

  Lys Asn Val Gln Gln Thr Val Ser Ala Lys Gly Pro Pro Glu Lys Arg
  225             230             235             240

  Met Arg Leu Gln
```

```
<210> 47
<211> 14
<212> DNA
<213> Homo sapiens

<400> 47
tttttttttt ttag                                              14
```

```
<210> 48
<211> 10
<212> DNA
<213> Homo sapiens

<400> 48
cttcaacctc                                                          10

<210> 49
<211> 496
<212> DNA
<213> Homo sapiens

<400> 49
gcaccatgta ccgagcactt cggctcctcg cgcgctcgcg tcccctcgtg cgggctccag 60
ccgcagcctt agcttcggct cccggcttgg gtggcgcggc cgtgccctcg ttttggcctc 120
cgaacgcggc tcgaatggca agccaaaatt ccttccggat agaatatgat acctttggtg 180
aactaaaggt gccaaatgat aagtattatg gcgcccagac cgtgagatct acgatgaact 240
ttaagattgg aggtgtgaca gaacgcatgc caaccccagt tattaaagct tttggcatct 300
tgaagcgagc ggccgctgaa gtaaaccagg attatggtct tgatccaaag attgctaatg 360
caataatgaa ggcagcagat gaggtagctg aaggtaaatt aaatgatcat tttcctctcg 420
tggtatggca gactggatca ggaactcaga caaatatgaa tgtaaatgaa gtcattagcc 480
aatagagcaa ttgaaa                                                  496

<210> 50
<211> 499
<212> DNA
<213> Homo sapiens

<400> 50
agaaaaagtc tatgtttgca gaaatacaga tccaagacaa agacaggatg ggcactgctg 60
gaaaagttat taaatgcaaa gcagctgtgc tttggggagca gaagcaaccc ttctccattg 120
aggaaataga agttgcccca ccaaagacta aagaagttcg cattaagatt ttggccacag 180
gaatctgtcg cacagatgac catgtgataa aaggaacaat ggtgtccaag tttccagtga 240
ttgtgggaca tgaggcaact gggattgtag agagcattgg agaaggagtg actacagtga 300
aaccaggtga caaagtcatc cctctctttc tgccacaatg tagagaatgc aatgcttgtc 360
gcaacccaga tggcaacctt tgcattagga gcgatattac tggtcgtgga gtactggctg 420
atggcaccac cagatttaca tgcaagggcg aaccagtcca ccacttcatg aacaccagta 480
catttaccga gtacacagt                                               499

<210> 51
<211> 887
<212> DNA
<213> Homo sapiens

<400> 51
gagtctgagc agaaaggaaa agcagccttg gcagccacgt tagaggaata caaagccaca 60
gtggccagtg accagataga gatgaatcgc ctgaaggctc agctggagaa tgaaaagcag 120
aaagtggcag agctgtattc tatccataac tctggagaca aatctgatat tcaggacctc 180
ctggagagtg tcaggctgga caaagaaaaa gcagagactt ggctagtag cttgcaggaa 240
gatctggctc atacccgaaa tgatgccaat cgattacagg atgccattgc taaggtagag 300
gatgaatacc gagccttcca agaagaagct aagaaacaaa ttgaagattt gaatatgacg 360
ttagaaaaat taagatcaga cctggatgaa aaagaaacag aaaggagtga catgaaagaa 420
accatctttg aacttgaaga tgaagtagaa caacatcgtg ctgtgaaact tcatgacaac 480
ctcattattt ctgatctaga gaatacagtt aaaaaactcc aggaccaaaa gcacgacatg 540
```

```
gaaagagaaa taaagacact ccacagaaga cttcgggaag aatctgcgga atggcggcag 600
tttcaggctg atctccagac tgcagtagtc attgcaaatg acattaaatc tgaagcccaa 660
gaggagattg gtgatctaaa gcgccggtta catgaggctc aagaaaaaaa tgagaaactc 720
acaaaagaat tggaggaaat aaagtcacgc aagcaagagg aggagcgagg cgggtataca 780
attacatgaa tgccgttgag agagatttgg cagccttaag gcagggaatg ggactgagta 840
gaaggtcctc gacttcctca gagccaactc ctacagtaaa aaccctc          887
```

```
<210> 52
<211> 491
<212> DNA
<213> Homo sapiens
```

```
<400> 52
ggcacgagct tttccaaaaa tcatgctgct cctttctcta aagttcttac attttatada 60
aaggaacctt tcactcttga ggcctactac agctctcctc aggatttgcc ctatccagat 120
cctgctatag ctcagttttc agttcagaaa gtcactcctc agtctgatgg ctccagttca 180
aaagtgaaag tcaaagttcg agtaaatgtc catggcattt tcagtgtgtc cagtgcatct 240
ttagtggagg ttcacaagtc tgaggaaaat gaggagccaa tggaaacaga tcagaatgca 300
aaggaggaag agaagatgca agtggaccag gaggaaccac atgttgaaga gcaacagcag 360
cagacaccag gcagaaaata aggcagagtc tgaagaaatg gagacctctc aagctggatc 420
caaggataaa aagatggacc aaccacccca agccaagaag gcaaaagtga agaccagtac 480
tgtggacctg g                                                 491
```

```
<210> 53
<211> 787
<212> DNA
<213> Homo sapiens
```

```
<400> 53
aagcagttga gtaggcagaa aaaagaacct cttcattaag gattaaaatg tataggccag 60
cacgtgtaac ttcgacttca agatttctga atccatatgt agtatgtttc attgtcgtcg 120
caggggtagt gatcctggca gtcaccatag ctctacttgt ttactttta gcttttgatc 180
aaaaatctta cttttatagg agcagttttc aactcctaaa tgttgaatat aatagtcagt 240
taaattcacc agctacacag gaatacagga ctttgagtgg aagaattgaa tctctgatta 300
ctaaaacatt caaagaatca aatttaagaa atcagttcat cagagctcat gttgccaaac 360
tgaggcaaga tggtagtggt gtgagagcgg atgttgtcat gaaattcaa ttcactagaa 420
ataacaatgg agcatcaatg aaaagcagaa ttgagtctgt tttacgacaa atgctgaata 480
actctggaaa cctggaaata aacccttcaa ctgagataac atcacttact gaccaggctg 540
cagcaaattg gcttattaat gaatgtgggg ccggtccaga cctaataaca ttgtctgagc 600
agagaatcct tggaggcact gaggctgagg agggaagctg gccgtggcaa gtcagtctgc 660
ggctcaataa tgcccaccac tgtggaggca gcctgatcaa taacatgtgg atcctgacag 720
cagctcactg cttcagaagc aactctaatc ctcgtgactg gattgccacg tctggtattt 780
ccacaac                                                      787
```

```
<210> 54
<211> 386
<212> DNA
<213> Homo sapiens
```

```
<400> 54
ggcattttca gtgtgtccag tgcatcttta gtggaggttc acaagtctga ggaaaatgag 60
gagccaatgg aaacagatca gaatgcaaag gaggaagaga agatgcaagt ggaccaggag 120
gaaccacatg ttgaagagca acagcagcag acaccagcag aaaataaggc agagtctgaa 180
gaaatggaga cctctcaagc tggatccaag gataaaaaga tggaccaacc accccaagcc 240
aagaaggcaa aagtgaagac cagtactgtg gacctgccaa tcgagaatca gctattatgg 300
```

cagatagaca gagagatgct caacttgtac attgaaaatg agggtaagat gatcatgcag 360
gataaactgg agaaggagcg gaatga   386

<210> 55
<211> 1462
<212> DNA
<213> Homo sapiens

<400> 55
aagcagttga gtaggcagaa aaaagaacct cttcattaag gattaaaatg tataggccag 60
cacgtgtaac ttcgacttca agatttctga atccatatgt agtatgtttc attgtcgtcg 120
caggggtagt gatcctggca gtcaccatag ctctacttgt ttactttta gcttttgatc 180
aaaaatctta ctttttatagg agcagtttc aactcctaaa tgttgaatat aatagtcagt 240
taaattcacc agctacacag gaatacagga ctttgagtgg aagaattgaa tctctgatta 300
ctaaacatt caaagaatca aatttaagaa atcagttcat cagagctcat gttgccaaac 360
tgaggcaaga tggtagtggt gtgagagcgg atgttgtcat gaaatttcaa ttcactagaa 420
ataacaatgg agcatcaatg aaaagcagaa ttgagtctgt tttacgacaa atgctgaata 480
actctggaaa cctggaaata aacccttcaa ctgagataac atcacttact gaccaggctg 540
cagcaaattg gcttattaat gaatgtgggg ccggtccaga cctaataaca ttgtctgagc 600
agagaatcct tggaggcact gaggctgagg agggaagctg gccgtggcaa gtcagtctgc 660
ggctcaataa tgcccaccac tgtggaggca gcctgatcaa taacatgtgg atcctgacag 720
cagctcactg cttcagaagc aactctaatc ctcgtgactg gattgccacg tctggtattt 780
ccacaacatt tcctaaacta agaatgagag taagaaatat tttaattcat aacaattata 840
aatctgcaac tcatgaaaat gacattgcac ttgtgagact tgagaacagt gtcacctta 900
ccaaagatat ccatagtgtg tgtctcccag ctgctaccca gaatattcca cctggctcta 960
ctgcttatgt aacaggatgg ggcgctcaag aatatgctgg ccacacagtt ccagagctaa 1020
ggcaaggaca ggtcagaata ataagtaatg atgtatgtaa tgcaccacat agttataatg 1080
gagccatctt gtctggaatg ctgtgtgctg gagtacctca aggtggagtg gacgcatgtc 1140
agggtgactc tggtggccca ctagtacaag aagactcacg gcggctttgg tttattgtgg 1200
ggatagtaag ctggggagat cagtgtggcc tgccggataa gccaggagtg tatactcgag 1260
tgacagcata cattgactgg attaggcaac aaactgggat ctagtgcaac aagtgcatcc 1320
ctgttgcaaa gtctgtatgc aggtgtgcct gtcttaaatt ccaaagcttt acatttcaac 1380
tgaaaaagaa actagaaatg tcctaattta acatcttgtt acataaatat ggtttaacaa 1440
aaaaaaaaa aaaaaactcg ag   1462

<210> 56
<211> 159
<212> PRT
<213> Homo sapiens

<400> 56
Thr Met Tyr Arg Ala Leu Arg Leu Leu Ala Arg Ser Arg Pro Leu Val
 1   5   10   15

Arg Ala Pro Ala Ala Ala Leu Ala Ser Ala Pro Gly Leu Gly Gly Ala
            20   25   30

Ala Val Pro Ser Phe Trp Pro Pro Asn Ala Ala Arg Met Ala Ser Gln
        35   40   45

Asn Ser Phe Arg Ile Glu Tyr Asp Thr Phe Gly Glu Leu Lys Val Pro
    50   55   60

Asn Asp Lys Tyr Tyr Gly Ala Gln Thr Val Arg Ser Thr Met Asn Phe
 65   70   75   80

Lys Ile Gly Gly Val Thr Glu Arg Met Pro Thr Pro Val Ile Lys Ala
              85   90   95

Phe Gly Ile Leu Lys Arg Ala Ala Ala Glu Val Asn Gln Asp Tyr Gly
              100   105   110

Leu Asp Pro Lys Ile Ala Asn Ala Ile Met Lys Ala Ala Asp Glu Val
          115   120   125

Ala Glu Gly Lys Leu Asn Asp His Phe Pro Leu Val Val Trp Gln Thr
      130   135   140

Gly Ser Gly Thr Gln Thr Asn Met Asn Val Asn Glu Val Ile Ser
145   150   155

<210> 57
<211> 165
<212> PRT
<213> Homo sapiens

<400> 57
Lys Lys Ser Met Phe Ala Glu Ile Gln Ile Gln Asp Lys Asp Arg Met
  1   5   10   15

Gly Thr Ala Gly Lys Val Ile Lys Cys Lys Ala Ala Val Leu Trp Glu
              20   25   30

Gln Lys Gln Pro Phe Ser Ile Glu Glu Ile Glu Val Ala Pro Pro Lys
          35   40   45

Thr Lys Glu Val Arg Ile Lys Ile Leu Ala Thr Gly Ile Cys Arg Thr
      50   55   60

Asp Asp His Val Ile Lys Gly Thr Met Val Ser Lys Phe Pro Val Ile
 65   70   75   80

Val Gly His Glu Ala Thr Gly Ile Val Glu Ser Ile Gly Glu Gly Val
              85   90   95

Thr Thr Val Lys Pro Gly Asp Lys Val Ile Pro Leu Phe Leu Pro Gln
          100   105   110

Cys Arg Glu Cys Asn Ala Cys Arg Asn Pro Asp Gly Asn Leu Cys Ile
          115   120   125

Arg Ser Asp Ile Thr Gly Arg Gly Val Leu Ala Asp Gly Thr Thr Arg
      130   135   140

Phe Thr Cys Lys Gly Glu Pro Val His His Phe Met Asn Thr Ser Thr
145   150   155   160

Phe Thr Glu Tyr Thr
              165

```
<210> 58
<211> 259
<212> PRT
<213> Homo sapiens

<400> 58
Glu Ser Glu Gln Lys Gly Lys Ala Ala Leu Ala Ala Thr Leu Glu Glu
  1               5                  10                  15

Tyr Lys Ala Thr Val Ala Ser Asp Gln Ile Glu Met Asn Arg Leu Lys
             20                  25                  30

Ala Gln Leu Glu Asn Glu Lys Gln Lys Val Ala Glu Leu Tyr Ser Ile
         35                  40                  45

His Asn Ser Gly Asp Lys Ser Asp Ile Gln Asp Leu Leu Glu Ser Val
     50                  55                  60

Arg Leu Asp Lys Glu Lys Ala Glu Thr Leu Ala Ser Ser Leu Gln Glu
 65                  70                  75                  80

Asp Leu Ala His Thr Arg Asn Asp Ala Asn Arg Leu Gln Asp Ala Ile
             85                  90                  95

Ala Lys Val Glu Asp Glu Tyr Arg Ala Phe Gln Glu Glu Ala Lys Lys
            100                 105                 110

Gln Ile Glu Asp Leu Asn Met Thr Leu Glu Lys Leu Arg Ser Asp Leu
        115                 120                 125

Asp Glu Lys Glu Thr Glu Arg Ser Asp Met Lys Glu Thr Ile Phe Glu
    130                 135                 140

Leu Glu Asp Glu Val Glu Gln His Arg Ala Val Lys Leu His Asp Asn
145                 150                 155                 160

Leu Ile Ile Ser Asp Leu Glu Asn Thr Val Lys Lys Leu Gln Asp Gln
                165                 170                 175

Lys His Asp Met Glu Arg Glu Ile Lys Thr Leu His Arg Arg Leu Arg
            180                 185                 190

Glu Glu Ser Ala Glu Trp Arg Gln Phe Gln Ala Asp Leu Gln Thr Ala
            195                 200                 205

Val Val Ile Ala Asn Asp Ile Lys Ser Glu Ala Gln Glu Glu Ile Gly
    210                 215                 220

Asp Leu Lys Arg Arg Leu His Glu Ala Gln Glu Lys Asn Glu Lys Leu
225                 230                 235                 240

Thr Lys Glu Leu Glu Glu Ile Lys Ser Arg Lys Gln Glu Glu Glu Arg
                245                 250                 255
```

Gly Gly Tyr

<210> 59
<211> 125
<212> PRT
<213> Homo sapiens

<400> 59
Gly Thr Ser Phe Ser Lys Asn His Ala Ala Pro Phe Ser Lys Val Leu
1   5   10   15

Thr Phe Tyr Arg Lys Glu Pro Phe Thr Leu Glu Ala Tyr Tyr Ser Ser
            20   25   30

Pro Gln Asp Leu Pro Tyr Pro Asp Pro Ala Ile Ala Gln Phe Ser Val
        35   40   45

Gln Lys Val Thr Pro Gln Ser Asp Gly Ser Ser Ser Lys Val Lys Val
        50   55   60

Lys Val Arg Val Asn Val His Gly Ile Phe Ser Val Ser Ser Ala Ser
65   70   75   80

Leu Val Glu Val His Lys Ser Glu Glu Asn Glu Glu Pro Met Glu Thr
                85   90   95

Asp Gln Asn Ala Lys Glu Glu Glu Lys Met Gln Val Asp Gln Glu Glu
            100   105   110

Pro His Val Glu Glu Gln Gln Gln Gln Thr Pro Gly Arg
            115   120   125

<210> 60
<211> 246
<212> PRT
<213> Homo sapiens

<400> 60
Met Tyr Arg Pro Ala Arg Val Thr Ser Thr Ser Arg Phe Leu Asn Pro
1   5   10   15

Tyr Val Val Cys Phe Ile Val Val Ala Gly Val Val Ile Leu Ala Val
            20   25   30

Thr Ile Ala Leu Leu Val Tyr Phe Leu Ala Phe Asp Gln Lys Ser Tyr
        35   40   45

Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln
    50   55   60

Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile
65   70   75   80

```
Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln
              85                  90                  95

Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val
          100                 105                 110

Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly
          115                 120                 125

Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met Leu Asn
      130                 135                 140

Asn Ser Gly Asn Leu Glu Ile Asn Pro Ser Thr Glu Ile Thr Ser Leu
145                 150                 155                 160

Thr Asp Gln Ala Ala Ala Asn Trp Leu Ile Asn Glu Cys Gly Ala Gly
              165                 170                 175

Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly Thr Glu
          180                 185                 190

Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu Asn Asn
          195                 200                 205

Ala His His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile Leu Thr
      210                 215                 220

Ala Ala His Cys Phe Arg Ser Asn Ser Asn Pro Arg Asp Trp Ile Ala
225                 230                 235                 240

Thr Ser Gly Ile Ser Thr
              245
```

<210> 61
<211> 128
<212> PRT
<213> Homo sapiens

<400> 61
```
Gly Ile Phe Ser Val Ser Ser Ala Ser Leu Val Glu Val His Lys Ser
 1           5                  10                  15

Glu Glu Asn Glu Glu Pro Met Glu Thr Asp Gln Asn Ala Lys Glu Glu
          20                  25                  30

Glu Lys Met Gln Val Asp Gln Glu Glu Pro His Val Glu Glu Gln Gln
      35                  40                  45

Gln Gln Thr Pro Ala Glu Asn Lys Ala Glu Ser Glu Glu Met Glu Thr
      50                  55                  60

Ser Gln Ala Gly Ser Lys Asp Lys Lys Met Asp Gln Pro Pro Gln Ala
65                  70                  75                  80

Lys Lys Ala Lys Val Lys Thr Ser Thr Val Asp Leu Pro Ile Glu Asn
```

```
                    85                 90                 95
Gln Leu Leu Trp Gln Ile Asp Arg Glu Met Leu Asn Leu Tyr Ile Glu
            100                 105                 110

Asn Glu Gly Lys Met Ile Met Gln Asp Lys Leu Glu Lys Glu Arg Asn
            115                 120                 125


<210> 62
<211> 418
<212> PRT
<213> Homo sapiens

<400> 62
Met Tyr Arg Pro Ala Arg Val Thr Ser Thr Ser Arg Phe Leu Asn Pro
  1             5                 10                 15

Tyr Val Val Cys Phe Ile Val Val Ala Gly Val Val Ile Leu Ala Val
            20                 25                 30

Thr Ile Ala Leu Leu Val Tyr Phe Leu Ala Phe Asp Gln Lys Ser Tyr
            35                 40                 45

Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln
  50                 55                 60

Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile
 65                 70                 75                 80

Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln
                85                 90                 95

Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val
            100                 105                 110

Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly
            115                 120                 125

Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met Leu Asn
            130                 135                 140

Asn Ser Gly Asn Leu Glu Ile Asn Pro Ser Thr Glu Ile Thr Ser Leu
            145                 150                 155                 160

Thr Asp Gln Ala Ala Ala Asn Trp Leu Ile Asn Glu Cys Gly Ala Gly
                165                 170                 175

Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly Thr Glu
            180                 185                 190

Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu Asn Asn
            195                 200                 205

Ala His His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile Leu Thr
```

```
            210                215                220

Ala Ala His Cys Phe Arg Ser Asn Ser Asn Pro Arg Asp Trp Ile Ala
225              230                235                    240

Thr Ser Gly Ile Ser Thr Thr Phe Pro Lys Leu Arg Met Arg Val Arg
             245                250                    255

Asn Ile Leu Ile His Asn Asn Tyr Lys Ser Ala Thr His Glu Asn Asp
             260                265                270

Ile Ala Leu Val Arg Leu Glu Asn Ser Val Thr Phe Thr Lys Asp Ile
         275                280                285

His Ser Val Cys Leu Pro Ala Ala Thr Gln Asn Ile Pro Pro Gly Ser
    290                295                300

Thr Ala Tyr Val Thr Gly Trp Gly Ala Gln Glu Tyr Ala Gly His Thr
305                310                315                320

Val Pro Glu Leu Arg Gln Gly Gln Val Arg Ile Ile Ser Asn Asp Val
             325                330                335

Cys Asn Ala Pro His Ser Tyr Asn Gly Ala Ile Leu Ser Gly Met Leu
         340                345                350

Cys Ala Gly Val Pro Gln Gly Gly Val Asp Ala Cys Gln Gly Asp Ser
         355                360                365

Gly Gly Pro Leu Val Gln Glu Asp Ser Arg Arg Leu Trp Phe Ile Val
370                375                380

Gly Ile Val Ser Trp Gly Asp Gln Cys Gly Leu Pro Asp Lys Pro Gly
385                390                395                400

Val Tyr Thr Arg Val Thr Ala Tyr Ile Asp Trp Ile Arg Gln Gln Thr
             405                410                415

Gly Ile
```

<210> 63
<211> 776
<212> DNA
<213> Homo sapiens

<400> 63
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cacagatggt gatagaggaa tccatcttgc agtcagataa agccctcact gatagagaga 60
aggcagtagc agtggatcgg gccaagaagg aggcagctga gaaggaacag gaacttttaa 120
aacagaaatt acaggagcag ccagcaacag atggaggctc aagataagag tcgcaaggaa 180
aactagccaa ctgaaggaga agctgcagat ggagagagaa cacctactga gagagcagat 240
tatgatgttg gagcacacgc agaaggtcca aaatgattgg cttcatgaag gatttaagaa 300
gaagtatgag gagatgaatg cagagataag tcaatttaaa cgtatgattg atactacaaa 360
aaatgatgat actccctgga ttgcacgaac cttggacaac cttgccgatg agctaactgc 420
aatattgtct gctcctgcta aattaattgg tcatggtgtc aaaggtgtga gctcactctt 480
```

taaaaagcat aagctcccct tttaaggata ttatagattg tacatatatg ctttggacta 540
tttttgatct gtatgttttt cattttcatt cagcaagttt tttttttttt tcagagtctt 600
actctgttgc ccaggctgga gtacagtggt gcaatctcag ctcactgcaa cctctgcctc 660
ctgggttcaa gagattcacc tgcctcagcc ccctagtagc tgggattata ggtgtacacc 720
accacaccca gctaattttt gtattttag tagagatggg gtttcactat gttggc   776

<210> 64
<211> 160
<212> DNA
<213> Homo sapiens

<400> 64
gcagcgctct cggttgcagt acccactgga aggacttagg cgctcgcgtg gacaccgcaa 60
gcccctcagt agcctcggcc caagaggcct gctttccact cgctagcccc gccggggtc 120
cgtgtcctgt ctcggtggcc ggacccgggc ccgagccga   160

<210> 65
<211> 72
<212> PRT
<213> Homo sapiens

<400> 65
Leu Ser Ala Met Gly Phe Thr Ala Ala Gly Ile Ala Ser Ser Ser Ile
1   5   10   15

Ala Ala Lys Met Met Ser Ala Ala Ala Ile Ala Asn Gly Gly Gly Val
            20   25   30

Ala Ser Gly Ser Leu Val Ala Thr Leu Gln Ser Leu Gly Ala Thr Gly
        35   40   45

Leu Ser Gly Leu Thr Lys Phe Ile Leu Gly Ser Ile Gly Ser Ala Ile
    50   55   60

Ala Ala Val Ile Ala Arg Phe Tyr
65   70

<210> 66
<211> 2581
<212> DNA
<213> Homo sapiens

<400> 66
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gctccgcagg gtgaggtggc tttgaccccg ggttgcccgg ccagcacgac cgaggaggtg 120
gctggacagc tggaggatga acgagaagc cgactgcccc acagacctgg aaatggccgc 180
ccccaaaggc caagaccgtt ggtcccagga agacatgctg actttgctgg aatgcatgaa 240
gaacaacctt ccatccaatg acagctccaa gttcaaaacc accgaatcac acatggactg 300
ggaaaaagta gcatttaaag acttttctgg agacatgtgc aagctcaaat gggtggagat 360
ttctaatgag gtgaggaagt tccgtacatt gacagaattg atcctcgatg ctcaggaaca 420
tgttaaaaat ccttacaaag gcaaaaaact caagaaacac ccagacttcc caagaagcc 480
cctgacccct tatttccgct tcttcatgga gaagcgggcc aagtatgcga aactccaccc 540
tgagatgagc aacctggacc taaccaagat tctgtccaag aaatacaagg agcttccgga 600
gaagaagaag atgaaatata ttcaggactt ccagagagag aaacaggagt tcgagcgaaa 660

```
cctggcccga ttcagggagg atcaccccga cctaatccag aatgccaaga aatcggacat  720
cccagagaag cccaaaaccc cccagcagct gtggtacacc cacgagaaga aggtgtatct  780
caaagtgcgg ccagatgcca ctacgaagga ggtgaaggac tccctgggga agcagtggtc  840
tcagctctcg gacaaaaaga ggctgaaatg gattcataag gccctggagc agcggaagga  900
gtacgaggag atcatgagag actatatcca gaagcaccca gagctgaaca tcagtgagga  960
gggtatcacc aagtccaccc tcaccaaggc cgaacgccag ctcaaggaca agtttgacgg 1020
gcgacccacc aagccacctc cgaacagcta ctcgctgtac tgcgcagagc tcatggccaa 1080
catgaaggac gtgcccagca cagagcgcat ggtgctgtgc agccagcagt ggaagctgct 1140
gtcccagaag gagaaggacg cctatcacaa gaagtgtgat cagaaaaaga aagattacga 1200
ggtggagctg ctccgtttcc tcgagagcct gcctgaggag gagcagcagc gggtcttggg 1260
ggaagagaag atgctgaaca tcaacaagaa gcaggccacc agccccgcct ccaagaagcc 1320
agcccaggaa gggggcaagg gcggctccga gaagcccaag cggcccgtgt cggccatgtt 1380
catcttctcg gaggagaaac ggcggcagct gcaggaggag cggcctgagc tctccgagag 1440
cgagctgacc cgcctgctgg cccgaatgtg gaacgacctg tctgagaaga agaaggccaa 1500
gtacaaggcc cgagaggcgg cgctcaaggc tcagtcggag aggaagcccg gcggggagcg 1560
cgaggaacgg ggcaagctgc ccgagtcccc caaaagagct gaggagatct ggcaacagag 1620
cgttatcggc gactacctgg cccgcttcaa gaatgaccgg gtgaaggcct tgaaagccat 1680
ggaaatgacc tggaataaca tggaaaagaa ggagaaactg atgtggatta agaaggcagc 1740
cgaagaccaa aagcgatatg agagagagct gagtgagatg cgggcacctc cagctgctac 1800
aaattcttcc aagaagatga aattccaggg agaacccaag aagcctccca tgaacggtta 1860
ccagaagttc tcccaggagc tgctgtccaa tggggagctg aaccacctgc cgctgaagga 1920
gcgcatggtg gagatcggca gtcgctggca gcgcatctcc cagagccaga aggagcacta 1980
caaaaagctg gccgaggagc agcaaaagca gtacaaggtg cacctggacc tctgggttaa 2040
gagcctgtct ccccaggacc gtgcagcata taaagagtac atctccaata aacgtaagag 2100
catgaccaag ctgcgaggcc caaaccccaa atccagccgg actactctgc agtccaagtc 2160
ggagtccgag gaggatgatg aagaggatga ggatgacgag gacgaggatg aagaagagga 2220
agatgatgag aatggggact cctctgaaga tggcggcgac tcctctgagt ccagcagcga 2280
ggacgagagc gaggatgggg atgagaatga agaggatgac gaggacgaag acgacgacga 2340
ggatgacgat gaggatgaag ataatgagtc cgagggcagc agctccagct cctcctcctt 2400
aggggactcc tcagactttg actccaactg aggcttagcc ccacccccagg ggagccaggg 2460
agagcccagg agctcccctc cccaactgac caccttgtt tcttccccat gttctgtccc 2520
ttgcccccct ggcctcccccc actttctttc tttctttaaa aaaaaaaaa aaaaactcga 2580
g                                                                  2581
```

<210> 67
<211> 764
<212> PRT
<213> Homo sapiens

<400> 67

```
Met Asn Gly Glu Ala Asp Cys Pro Thr Asp Leu Glu Met Ala Ala Pro
1               5                   10                  15

Lys Gly Gln Asp Arg Trp Ser Gln Glu Asp Met Leu Thr Leu Leu Glu
            20                  25                  30

Cys Met Lys Asn Asn Leu Pro Ser Asn Asp Ser Ser Lys Phe Lys Thr
        35                  40                  45

Thr Glu Ser His Met Asp Trp Glu Lys Val Ala Phe Lys Asp Phe Ser
    50                  55                  60

Gly Asp Met Cys Lys Leu Lys Trp Val Glu Ile Ser Asn Glu Val Arg
65                  70                  75                  80
```

```
Lys Phe Arg Thr Leu Thr Glu Leu Ile Leu Asp Ala Gln Glu His Val
              85                    90                    95

Lys Asn Pro Tyr Lys Gly Lys Lys Leu Lys Lys His Pro Asp Phe Pro
            100                   105                   110

Lys Lys Pro Leu Thr Pro Tyr Phe Arg Phe Phe Met Glu Lys Arg Ala
            115                   120                   125

Lys Tyr Ala Lys Leu His Pro Glu Met Ser Asn Leu Asp Leu Thr Lys
        130                   135                   140

Ile Leu Ser Lys Lys Tyr Lys Glu Leu Pro Glu Lys Lys Lys Met Lys
145                   150                   155                   160

Tyr Ile Gln Asp Phe Gln Arg Glu Lys Gln Glu Phe Glu Arg Asn Leu
                165                   170                   175

Ala Arg Phe Arg Glu Asp His Pro Asp Leu Ile Gln Asn Ala Lys Lys
            180                   185                   190

Ser Asp Ile Pro Glu Lys Pro Lys Thr Pro Gln Gln Leu Trp Tyr Thr
            195                   200                   205

His Glu Lys Lys Val Tyr Leu Lys Val Arg Pro Asp Ala Thr Thr Lys
    210                   215                   220

Glu Val Lys Asp Ser Leu Gly Lys Gln Trp Ser Gln Leu Ser Asp Lys
225                   230                   235                   240

Lys Arg Leu Lys Trp Ile His Lys Ala Leu Glu Gln Arg Lys Glu Tyr
            245                   250                   255

Glu Glu Ile Met Arg Asp Tyr Ile Gln Lys His Pro Glu Leu Asn Ile
            260                   265                   270

Ser Glu Glu Gly Ile Thr Lys Ser Thr Leu Thr Lys Ala Glu Arg Gln
    275                   280                   285

Leu Lys Asp Lys Phe Asp Gly Arg Pro Thr Lys Pro Pro Pro Asn Ser
    290                   295                   300

Tyr Ser Leu Tyr Cys Ala Glu Leu Met Ala Asn Met Lys Asp Val Pro
305                   310                   315                   320

Ser Thr Glu Arg Met Val Leu Cys Ser Gln Gln Trp Lys Leu Leu Ser
            325                   330                   335

Gln Lys Glu Lys Asp Ala Tyr His Lys Lys Cys Asp Gln Lys Lys Lys
        340                   345                   350

Asp Tyr Glu Val Glu Leu Leu Arg Phe Leu Glu Ser Leu Pro Glu Glu
        355                   360                   365

Glu Gln Gln Arg Val Leu Gly Glu Glu Lys Met Leu Asn Ile Asn Lys
```

```
        370                375                 380
Lys Gln Ala Thr Ser Pro Ala Ser Lys Lys Pro Ala Gln Glu Gly Gly
385                390                  395                   400

 Lys Gly Gly Ser Glu Lys Pro Lys Arg Pro Val Ser Ala Met Phe Ile
              405              410                    415

 Phe Ser Glu Glu Lys Arg Arg Gln Leu Gln Glu Glu Arg Pro Glu Leu
              420              425                   430

 Ser Glu Ser Glu Leu Thr Arg Leu Leu Ala Arg Met Trp Asn Asp Leu
        435               440               445

 Ser Glu Lys Lys Lys Ala Lys Tyr Lys Ala Arg Glu Ala Ala Leu Lys
    450               455               460

Ala Gln Ser Glu Arg Lys Pro Gly Gly Glu Arg Glu Glu Arg Gly Lys
465               470               475               480

Leu Pro Glu Ser Pro Lys Arg Ala Glu Glu Ile Trp Gln Gln Ser Val
              485               490               495

Ile Gly Asp Tyr Leu Ala Arg Phe Lys Asn Asp Arg Val Lys Ala Leu
              500               505               510

Lys Ala Met Glu Met Thr Trp Asn Asn Met Glu Lys Lys Glu Lys Leu
        515               520               525

Met Trp Ile Lys Lys Ala Ala Glu Asp Gln Lys Arg Tyr Glu Arg Glu
    530               535               540

Leu Ser Glu Met Arg Ala Pro Pro Ala Ala Thr Asn Ser Ser Lys Lys
545               550               555               560

Met Lys Phe Gln Gly Glu Pro Lys Lys Pro Pro Met Asn Gly Tyr Gln
              565               570               575

Lys Phe Ser Gln Glu Leu Leu Ser Asn Gly Glu Leu Asn His Leu Pro
              580               585               590

Leu Lys Glu Arg Met Val Glu Ile Gly Ser Arg Trp Gln Arg Ile Ser
         595               600               605

Gln Ser Gln Lys Glu His Tyr Lys Lys Leu Ala Glu Glu Gln Gln Lys
    610               615               620

Gln Tyr Lys Val His Leu Asp Leu Trp Val Lys Ser Leu Ser Pro Gln
625               630               635               640

Asp Arg Ala Ala Tyr Lys Glu Tyr Ile Ser Asn Lys Arg Lys Ser Met
              645               650               655

Thr Lys Leu Arg Gly Pro Asn Pro Lys Ser Ser Arg Thr Thr Leu Gln
              660               665               670
```

```
Ser Lys Ser Glu Ser Glu Glu Asp Asp Glu Glu Asp Glu Asp Asp Glu
    675             680             685

Asp Glu Asp Glu Glu Glu Glu Asp Asp Glu Asn Gly Asp Ser Ser Glu
    690             695             700

Asp Gly Gly Asp Ser Ser Glu Ser Ser Ser Glu Asp Glu Ser Glu Asp
705             710             715             720

Gly Asp Glu Asn Glu Glu Asp Asp Glu Asp Glu Asp Asp Asp Glu Asp
            725             730             735

Asp Asp Glu Asp Glu Asp Asn Glu Ser Glu Gly Ser Ser Ser Ser Ser
        740             745             750

Ser Ser Leu Gly Asp Ser Ser Asp Phe Asp Ser Asn
    755             760
```

<210> 68
<211> 434
<212> DNA
<213> Homo sapiens

<400> 68
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ctaagatgct ggatgctgaa gacatcgtcg gaactgcccg gccagatgag aaagccatta 60
tgacttatgt gtctagcttc tatcatgcct tctctggagc ccagaaggca gaaacagcag 120
ccaatcgcat ctgcaaagtg ttggcggtca atcaagagaa cgagcagctt atggaagact 180
atgagaagct ggccagtgat ctgttggagt ggatccgccg caccatccca tggctggaga 240
atcgggtgcc tgagaacacc atgcatgcca tgcagcagaa gctggaggac ttccgagact 300
atagacgcct gcacaagccg cccaaggtgc aggagaagtg ccagctggag atcaacttta 360
acacgctgca gaccaaactg cggctcagca accggcctgc cttcatgccc tccgagggca 420
ggatggtctc ggat                                                  434
```

<210> 69
<211> 244
<212> DNA
<213> Homo sapiens

<400> 69
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aggcagcatg ctcgttgaga gtcatcacca ctccctaatc tcaagtacgc agggacacaa 60
acactgcgga aggccgcagg gtcctctgcc taggaaaacc agagaccttt gttcacttgt 120
ttatgtgctg accttccctc cactattgtc ctgtgaccct gccaaatccc cctttgtgag 180
aaacacccaa gaatgatcaa taaaaaataa attaatttag gaaaaaaaaa aaaaaaaact 240
cgag                                                             244
```

<210> 70
<211> 437
<212> DNA
<213> Homo sapiens

<400> 70
```
ctgggacggg agcgtccagc gggactcgaa ccccagatgt gaaggcgttt ctggaaagtc 60
cttggtccct ggatccagcg tcggccagcc cagagcccgt gccgcacatc cttgcgtcct 120
```

```
ccaggcagtg ggaccccgcg agctgcacgt ccctgggcac ggacaagtgt gaggcactgt 180
tggggctgtg ccaggtgcgg ggtgggctgc cccctttctc agaaccttcc agcctggtgc 240
cgtggccccc aggccggagt cttcctaagg ctgtgaggcc acccctgtcc tggcctccgt 300
tctcgcagca gcagaccttg cccgtgatga gcggggaggc ccttggctgg ctgggccagg 360
ctggttccct ggccatgggg gctgcacctc tgggggagcc agccaaggag gaccccatgc 420
tggcgcagga agccggg                                                 437
```

<210> 71
<211> 271
<212> DNA
<213> Homo sapiens

<400> 71
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gcgcagagtt ctgtcgtcca ccatcgagtg aggaagagag cattggttcc cctgagatag 60
aagagatggc tctcttcagt gcccagtctc catacattaa cccgatcatc ccctttactg 120
gaccaatcca aggagggctg caggagggac ttcaggtgac cctccagggg actaccgaga 180
gttttgcaca aagtttgtg gtgaactttt cagaacagct tcaatggaga tgacttggcc 240
ttccacttca accccggtta tgaggaagga g.                                 271
```

<210> 72
<211> 290
<212> DNA
<213> Homo sapiens

<400> 72
```
ccgagcccta cccggaggtc tccagaatcc ccaccgtcag gggatgcaac ggctccctgt 60
ctggtgccct ctcctgctgc gaggactcgg cccagggctc gggcccgccc aaggcccta 120
cggtggccga gggtcccagc tcctgccttc ggcggaacgt gatcagcgag agggagcgca 180
ggaagcggat gtcgttgagc tgtgagcgtc tgcgggccct gctgccccag ttcgatggcc 240
ggcgggagga catggcctcg gtcctggaga tgtctgttgc aattcctgcg             290
```

<210> 73
<211> 144
<212> PRT
<213> Homo sapiens

<400> 73
```
Lys Met Leu Asp Ala Glu Asp Ile Val Gly Thr Ala Arg Pro Asp Glu
 1               5                  10                  15

Lys Ala Ile Met Thr Tyr Val Ser Ser Phe Tyr His Ala Phe Ser Gly
            20                  25                  30

Ala Gln Lys Ala Glu Thr Ala Ala Asn Arg Ile Cys Lys Val Leu Ala
        35                  40                  45

Val Asn Gln Glu Asn Glu Gln Leu Met Glu Asp Tyr Glu Lys Leu Ala
    50                  55                  60

Ser Asp Leu Leu Glu Trp Ile Arg Arg Thr Ile Pro Trp Leu Glu Asn
65                  70                  75                  80

Arg Val Pro Glu Asn Thr Met His Ala Met Gln Gln Lys Leu Glu Asp
                85                  90                  95
```

Phe Arg Asp Tyr Arg Arg Leu His Lys Pro Pro Lys Val Gln Glu Lys
         100   105   110

Cys Gln Leu Glu Ile Asn Phe Asn Thr Leu Gln Thr Lys Leu Arg Leu
         115   120   125

Ser Asn Arg Pro Ala Phe Met Pro Ser Glu Gly Arg Met Val Ser Asp
         130   135   140

<210> 74
<211> 64
<212> PRT
<213> Homo sapiens

<400> 74
Gly Ser Met Leu Val Glu Ser His His His Ser Leu Ile Ser Ser Thr
  1   5   10   15

Gln Gly His Lys His Cys Gly Arg Pro Gln Gly Pro Leu Pro Arg Lys
         20   25   30

Thr Arg Asp Leu Cys Ser Leu Val Tyr Val Leu Thr Phe Pro Pro Leu
         35   40   45

Leu Ser Cys Asp Pro Ala Lys Ser Pro Phe Val Arg Asn Thr Gln Glu
     50   55   60

<210> 75
<211> 145
<212> PRT
<213> Homo sapiens

<400> 75
Gly Thr Gly Ala Ser Ser Gly Thr Arg Thr Pro Asp Val Lys Ala Phe
  1   5   10   15

Leu Glu Ser Pro Trp Ser Leu Asp Pro Ala Ser Ala Ser Pro Glu Pro
         20   25   30

Val Pro His Ile Leu Ala Ser Ser Arg Gln Trp Asp Pro Ala Ser Cys
         35   40   45

Thr Ser Leu Gly Thr Asp Lys Cys Glu Ala Leu Leu Gly Leu Cys Gln
     50   55   60

Val Arg Gly Gly Leu Pro Pro Phe Ser Glu Pro Ser Ser Leu Val Pro
 65   70   75   80

Trp Pro Pro Gly Arg Ser Leu Pro Lys Ala Val Arg Pro Pro Leu Ser
                 85   90   95

Trp Pro Pro Phe Ser Gln Gln Gln Thr Leu Pro Val Met Ser Gly Glu
             100   105   110

Ala Leu Gly Trp Leu Gly Gln Ala Gly Ser Leu Ala Met Gly Ala Ala
        115   120   125

Pro Leu Gly Glu Pro Ala Lys Glu Asp Pro Met Leu Ala Gln Glu Ala
    130   135   140

Gly
145

<210> 76
<211> 69
<212> PRT
<213> Homo sapiens

<400> 76
Ala Glu Phe Cys Arg Pro Pro Ser Ser Glu Glu Glu Ser Ile Gly Ser
  1   5   10   15

Pro Glu Ile Glu Glu Met Ala Leu Phe Ser Ala Gln Ser Pro Tyr Ile
              20   25   30

Asn Pro Ile Ile Pro Phe Thr Gly Pro Ile Gln Gly Gly Leu Gln Glu
        35   40   45

Gly Leu Gln Val Thr Leu Gln Gly Thr Thr Glu Ser Phe Ala Gln Lys
     50   55   60

Phe Val Val Asn Phe
  65

<210> 77
<211> 96
<212> PRT
<213> Homo sapiens

<400> 77
Glu Pro Tyr Pro Glu Val Ser Arg Ile Pro Thr Val Arg Gly Cys Asn
  1   5   10   15

Gly Ser Leu Ser Gly Ala Leu Ser Cys Cys Glu Asp Ser Ala Gln Gly
            20   25   30

Ser Gly Pro Pro Lys Ala Pro Thr Val Ala Glu Gly Pro Ser Ser Cys
         35   40   45

Leu Arg Arg Asn Val Ile Ser Glu Arg Glu Arg Arg Lys Arg Met Ser
     50   55   60

Leu Ser Cys Glu Arg Leu Arg Ala Leu Leu Pro Gln Phe Asp Gly Arg
  65   70   75   80

Arg Glu Asp Met Ala Ser Val Leu Glu Met Ser Val Ala Ile Pro Ala

85   90   95

<210> 78
<211> 2076
<212> DNA
<213> Homo sapiens

<400> 78
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gaatctgtcg cacagatgac catgtgataa aaggaacaat ggtgtccaag tttccagtga 240
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aaccaggtga caaagtcatc cctctctttc tgccacaatg tagagaatgc aatgcttgtc 360
gcaacccaga tggcaacctt tgcattagga gcgatattac tggtcgtgga gtactggctg 420
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catttaccga gtacacagtg gtggatgaat cttctgttgc taagattgat gatgcagctc 540
ctcctgagaa agtctgttta attggctgtg ggttttccac tggatatggc gctgctgtta 600
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tagaatcata ttatcatact tatcataatg ttcaatttga tacagtagaa ttgcaagtcc 1920
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<210> 79
<211> 2790
<212> DNA
<213> Homo sapiens

<400> 79
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cagggggtagt gatcctggca gtcaccatag ctctacttgt ttactttta gcttttgatc 180
aaaaatctta ctttttatagg agcagttttc aactcctaaa tgttgaatat aatagtcagt 240

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taaattcacc agctacacag gaatacagga ctttgagtgg aagaattgaa tctctgatta 300
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ataacaatgg agcatcaatg aaaagcagaa ttgagtctgt tttacgacaa atgctgaata 480
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agatattttt catgatgacc catgaaaaat atcactcatc tacataaagg agagactata 2040
tctattttat agagaagcta agaaatatac ctacacaaac ttgtcaggtg ctttacaact 2100
acatagtact ttttaacaac aaaataataa ttttaagaat gaaaaattta atcatcggga 2160
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tcttggcgaa ctgtacaaac aaatctttgc tatactttat ttcaaataaa ttcttttttga 2760
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<210> 80
<211> 1460
<212> DNA
<213> Homo sapiens

<400> 80
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gtcgcagggg tagtgatcct ggcagtcacc atagctctac ttgtttactt tttagctttt 180
gatcaaaaat cttactttta taggagcagt tttcaactcc taaatgttga atataatagt 240
cagttaaatt caccagctac acaggaatac aggactttga gtggaagaat tgaatctctg 300
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attactaaaa cattcaaaga atcaaattta agaaatcagt tcatcagagc tcatgttgcc 360
aaactgaggc aagatggtag tggtgtgaga gcggatgttg tcatgaaatt tcaattcact 420
agaaataaca atggagcatc aatgaaaagc agaattgagt ctgtttacg  acaaatgctg 480
aataactctg gaaacctgga aataaaccct tcaactgaga taacatcact tactgaccag 540
gctgcagcaa attggcttat taatgaatgt ggggccggtc cagacctaat aacattgtct 600
gagcagagaa tccttggagg cactgaggct gaggagggaa gctggccgtg gcaagtcagt 660
ctgcggctca ataatgccca ccactgtgga ggcagcctga tcaataacat gtggatcctg 720
acagcagctc actgcttcag aagcaactct aatcctcgtg actggattgc cacgtctggt 780
atttccacaa catttcctaa actaagaatg agagtaagaa atattttaat tcataacaat 840
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atccctgttg caaagtctgt atgcaggtgt gcctgtctta aattccaaag ctttacattt 1380
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acaaaaaaaa aaaaaaaaaa                                            1460
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<210> 81

<211> 386

<212> PRT

<213> Homo sapiens

<400> 81

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Met Phe Ala Glu Ile Gln Ile Gln Asp Lys Asp Arg Met Gly Thr Ala
1               5                   10                  15

Gly Lys Val Ile Lys Cys Lys Ala Ala Val Leu Trp Glu Gln Lys Gln
            20                  25                  30

Pro Phe Ser Ile Glu Glu Ile Glu Val Ala Pro Pro Lys Thr Lys Glu
        35                  40                  45

Val Arg Ile Lys Ile Leu Ala Thr Gly Ile Cys Arg Thr Asp Asp His
    50                  55                  60

Val Ile Lys Gly Thr Met Val Ser Lys Phe Pro Val Ile Val Gly His
65                  70                  75                  80

Glu Ala Thr Gly Ile Val Glu Ser Ile Gly Glu Gly Val Thr Thr Val
                85                  90                  95

Lys Pro Gly Asp Lys Val Ile Pro Leu Phe Leu Pro Gln Cys Arg Glu
            100                 105                 110

Cys Asn Ala Cys Arg Asn Pro Asp Gly Asn Leu Cys Ile Arg Ser Asp
            115                 120                 125

Ile Thr Gly Arg Gly Val Leu Ala Asp Gly Thr Thr Arg Phe Thr Cys
        130                 135                 140

Lys Gly Lys Pro Val His His Phe Met Asn Thr Ser Thr Phe Thr Glu
```

```
      145                 150                 155                 160

      Tyr Thr Val Val Asp Glu Ser Ser Val Ala Lys Ile Asp Asp Ala Ala
                      165                 170                 175

      Pro Pro Glu Lys Val Cys Leu Ile Gly Cys Gly Phe Ser Thr Gly Tyr
                      180                 185                 190

      Gly Ala Ala Val Lys Thr Gly Lys Val Lys Pro Gly Ser Thr Cys Val
                      195                 200                 205

      Val Phe Gly Leu Arg Gly Val Gly Leu Ser Val Ile Met Gly Cys Lys
          210                 215                 220

      Ser Ala Gly Ala Ser Arg Ile Ile Gly Ile Asp Leu Asn Lys Asp Lys
      225                 230                 235                 240

      Phe Glu Lys Ala Met Ala Val Gly Ala Thr Glu Cys Ile Ser Pro Lys
                      245                 250                 255

      Asp Ser Thr Lys Pro Ile Ser Glu Val Leu Ser Glu Met Thr Gly Asn
                      260                 265                 270

      Asn Val Gly Tyr Thr Phe Glu Val Ile Gly His Leu Glu Thr Met Ile
                      275                 280                 285

      Asp Ala Leu Ala Ser Cys His Met Asn Tyr Gly Thr Ser Val Val Val
          290                 295                 300

      Gly Val Pro Pro Ser Ala Lys Met Leu Thr Tyr Asp Pro Met Leu Leu
      305                 310                 315                 320

      Phe Thr Gly Arg Thr Trp Lys Gly Cys Val Phe Gly Gly Leu Lys Ser
                      325                 330                 335

      Arg Asp Asp Val Pro Lys Leu Val Thr Glu Phe Leu Ala Lys Lys Phe
                      340                 345                 350

      Asp Leu Asp Gln Leu Ile Thr His Val Leu Pro Phe Lys Lys Ile Ser
                      355                 360                 365

      Glu Gly Phe Glu Leu Leu Asn Ser Gly Gln Ser Ile Arg Thr Val Leu
          370                 375                 380

      Thr Phe
      385
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<210> 82
<211> 418
<212> PRT
<213> Homo sapiens

<400> 82
Met Tyr Arg Pro Ala Arg Val Thr Ser Thr Ser Arg Phe Leu Asn Pro
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| | | | 1 | | | | | 5 | | | | | 10 | | | | | 15 |

Tyr Val Val Cys Phe Ile Val Val Ala Gly Val Val Ile Leu Ala Val
            20   25   30

Thr Ile Ala Leu Leu Val Tyr Phe Leu Ala Phe Asp Gln Lys Ser Tyr
        35   40   45

Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln
    50   55   60

Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile
65   70   75   80

Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln
            85   90   95

Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val
            100   105   110

Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly
        115   120   125

Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met Leu Asn
130   135   140

Asn Ser Gly Asn Leu Glu Ile Asn Pro Ser Thr Glu Ile Thr Ser Leu
145   150   155   160

Thr Asp Gln Ala Ala Ala Asn Trp Leu Ile Asn Glu Cys Gly Ala Gly
            165   170   175

Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly Thr Glu
            180   185   190

Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu Asn Asn
            195   200   205

Ala His His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile Leu Thr
    210   215   220

Ala Ala His Cys Phe Arg Ser Asn Ser Asn Pro Arg Asp Trp Ile Ala
225   230   235   240

Thr Ser Gly Ile Ser Thr Thr Phe Pro Lys Leu Arg Met Arg Val Arg
            245   250   255

Asn Ile Leu Ile His Asn Asn Tyr Lys Ser Ala Thr His Glu Asn Asp
            260   265   270

Ile Ala Leu Val Arg Leu Glu Asn Ser Val Thr Phe Thr Lys Asp Ile
            275   280   285

His Ser Val Cys Leu Pro Ala Ala Thr Gln Asn Ile Pro Pro Gly Ser
    290   295   300

```
Thr Ala Tyr Val Thr Gly Trp Gly Ala Gln Glu Tyr Ala Gly His Thr
305              310              315              320

Val Pro Glu Leu Arg Gln Gly Gln Val Arg Ile Ile Ser Asn Asp Val
              325              330              335

Cys Asn Ala Pro His Ser Tyr Asn Gly Ala Ile Leu Ser Gly Met Leu
              340              345              350

Cys Ala Gly Val Pro Gln Gly Gly Val Asp Ala Cys Gln Gly Asp Ser
     355              360              365

Gly Gly Pro Leu Val Gln Glu Asp Ser Arg Arg Leu Trp Phe Ile Val
   370              375              380

Gly Ile Val Ser Trp Gly Asp Gln Cys Gly Leu Pro Asp Lys Pro Gly
385              390              395              400

Val Tyr Thr Arg Val Thr Ala Tyr Leu Asp Trp Ile Arg Gln Gln Thr
              405              410              415

Gly Ile
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<210> 83
<211> 418
<212> PRT
<213> Homo sapiens

<400> 83
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Met Tyr Arg Pro Ala Arg Val Thr Ser Thr Ser Arg Phe Leu Asn Pro
1              5              10              15

Tyr Val Val Cys Phe Ile Val Val Ala Gly Val Val Ile Leu Ala Val
              20              25              30

Thr Ile Ala Leu Leu Val Tyr Phe Leu Ala Phe Asp Gln Lys Ser Tyr
     35              40              45

Phe Tyr Arg Ser Ser Phe Gln Leu Leu Asn Val Glu Tyr Asn Ser Gln
   50              55              60

Leu Asn Ser Pro Ala Thr Gln Glu Tyr Arg Thr Leu Ser Gly Arg Ile
65              70              75              80

Glu Ser Leu Ile Thr Lys Thr Phe Lys Glu Ser Asn Leu Arg Asn Gln
              85              90              95

Phe Ile Arg Ala His Val Ala Lys Leu Arg Gln Asp Gly Ser Gly Val
          100              105              110

Arg Ala Asp Val Val Met Lys Phe Gln Phe Thr Arg Asn Asn Asn Gly
          115              120              125
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```
Ala Ser Met Lys Ser Arg Ile Glu Ser Val Leu Arg Gln Met Leu Asn
    130             135             140

Asn Ser Gly Asn Leu Glu Ile Asn Pro Ser Thr Glu Ile Thr Ser Leu
145             150             155             160

Thr Asp Gln Ala Ala Ala Asn Trp Leu Ile Asn Glu Cys Gly Ala Gly
                165             170             175

Pro Asp Leu Ile Thr Leu Ser Glu Gln Arg Ile Leu Gly Gly Thr Glu
            180             185             190

Ala Glu Glu Gly Ser Trp Pro Trp Gln Val Ser Leu Arg Leu Asn Asn
        195             200             205

Ala His His Cys Gly Gly Ser Leu Ile Asn Asn Met Trp Ile Leu Thr
    210             215             220

Ala Ala His Cys Phe Arg Ser Asn Ser Asn Pro Arg Asp Trp Ile Ala
225             230             235             240

Thr Ser Gly Ile Ser Thr Thr Phe Pro Lys Leu Arg Met Arg Val Arg
                245             250             255

Asn Ile Leu Ile His Asn Asn Tyr Lys Ser Ala Thr His Glu Asn Asp
        260             265             270

Ile Ala Leu Val Arg Leu Glu Asn Ser Val Thr Phe Thr Lys Asp Ile
    275             280             285

His Ser Val Cys Leu Pro Ala Ala Thr Gln Asn Ile Pro Pro Gly Ser
    290             295             300

Thr Ala Tyr Val Thr Gly Trp Gly Ala Gln Glu Tyr Ala Gly His Thr
305             310             315             320

Val Pro Glu Leu Arg Gln Gly Gln Val Arg Ile Ile Ser Asn Asp Val
            325             330             335

Cys Asn Ala Pro His Ser Tyr Asn Gly Ala Ile Leu Ser Gly Met Leu
            340             345             350

Cys Ala Gly Val Pro Gln Gly Gly Val Asp Ala Cys Gln Gly Asp Ser
    355             360             365

Gly Gly Pro Leu Val Gln Glu Asp Ser Arg Arg Leu Trp Phe Ile Val
    370             375             380

Gly Ile Val Ser Trp Gly Asp Gln Cys Gly Leu Pro Asp Lys Pro Gly
385             390             395             400

Val Tyr Thr Arg Val Thr Ala Tyr Leu Asp Trp Ile Arg Gln Gln Thr
            405             410             415
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Gly Ile

<210> 84
<211> 489
<212> DNA
<213> Homo sapiens

<400> 84
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aattacagag gagtttcatg gcactaagtc aagatattca gaaaacaata aagaagacag 180
cacgtcggga gcagcttatg agagaagaag ctgaacagaa acgtttaaaa actgtacttg 240
agctacagta tgttttggac aaattgggag atgatgaagt gcggactgac ctgaaacaag 300
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ccattcacct gtgggacctg ctggaaggga aggaaaaacc tgtatgtgga accacctata 480
aagttctaa   489

<210> 85
<211> 304
<212> DNA
<213> Homo sapiens

<400> 85
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agctggagaa ggagaagagc gagatgaaga tggagatcga tgacctcgct tgtaacatgg 180
aggtcatctc caaatctaag ggaaaccttg agaagatgtg ccgcacactg gaggaccaag 240
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agag   304

<210> 86
<211> 296
<212> DNA
<213> Homo sapiens

<400> 86
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ttccttaagg attaaaatgt ttagggcaac acgtgttact tccacttcca gatttctgaa 120
tccatatgtt gtatgtttcc ttgtcctccc aggggttgtg atcctggcag tccccatagc 180
tctacttgtt tacttttag cttttgatca aaaatcttac ttttattgga gcaattttcc 240
actcccaaat gttgaatata atagtccgtt taattccccc gcttcaccgg gaattc 296

<210> 87
<211> 904
<212> DNA
<213> Homo sapiens

<400> 87
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tatctgatcg ttctaaaaaa gagttgtccc cggttttaac cagtgaagtt catagtgttc 180
gtgcaggacg gcatcttgct accaaaattga atattttagt acagcaacat tttgacttgg 240
cttcaactac tattacaaat attccaatga aggaagaaca gcatgctaac acatctgcca 300
attatgatgt ggagctactt catcacaaag atgcacatgt agatttcctg aaaagtggtg 360

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cacctgtaga tgtaaatagt agaccttcct cctgccttac taattttctt ctaaatggtc 540
gttctgtttt attggaacaa ccacgaaagt caggttctaa agtcattagt catatgctta 600
gtagccatgg aggagagatt tttttgcacg tccttagcag ttctcgatcc attctagaag 660
atccaccttc aattagtgaa ggatgtggag gaagagttac agactaccgg attacagatt 720
ttggtgaatt tatgagggga aaacagatta actccttttc tacaccccag atataaaatc 780
gatggaagtc ttgaggtccc tttggaaccg agccaaaaga tcagttaaaa aaacataccc 840
gttactggcc tatgatttca aaaacccacc atttttaaca tgcaagcggt agttccgtta 900
acca                                                          904


<210> 88
<211> 387
<212> DNA
<213> Homo sapiens


<400> 88
cgtctctccc ccagtttgcc gttcacccgg agcgctcggg acttgccgat agtggtgacg 60
gcggcaacat gtctgtggct ttcgcggccc cgaggcagcg aggcaagggg gagatcactc 120
ccgctgcgat tcagaagatg ttggatgaca ataaccatct tattcagtgt ataatggact 180
ctcagaataa aggaaagacc tcagagtgtt ctcagtatca gcagatgttg cacacaaact 240
tggtatacct tgctacaata gcagattcta atcaaaatat gcagtctctt ttaccagcac 300
cacccacaca gaatatgcct atgggtcctg gagggatgaa tcagagcggg cctcccccac 360
ctccacgctc tcacaacatg ccttcaa                                 387


<210> 89
<211> 481
<212> DNA
<213> Homo sapiens


<400> 89
tgttcttgga cctgcggtgc tatagagcag gctcttctag gttggcagtt gccatggaat 60
ctggacccaa aatgttggcc cccgtttgcc tggtggaaaa taacaatgag cagctattgg 120
tgaaccagca agctatacag attcttgaaa agatttctca gccagtggtg gtggtggcca 180
ttgtaggact gtaccgtaca gggaaatcct acttgatgaa ccatctggca ggacagaatc 240
atggcttccc tctgggctcc acggtgcagt ctgaaaccaa gggcatctgg atgtggtgcg 300
tgccccaccc atccaagcca aaccacaccc tggtccttct ggacaccgaa ggtctgggcg 360
atgtggaaaa gggtgaccct aagaatgact cctggatctt tgccctggct gtgctcctgt 420
gcagcacctt tgtctacaac agcatgagca ccatcaacca ccaggccctg gagcagctgc 480
a                                                             481


<210> 90
<211> 491
<212> DNA
<213> Homo sapiens


<400> 90
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accagcaagc tatacagatt cttgaaaaga ttctcagcc agtggtggtg gtggccattg 180
taggactgta ccgtacaggg aaatcctact tgatgaacca tctggcagga cagaatcatg 240
gcttccctct gggctccacg gtgcagtctg aaaccaaggg catctggatg tggtgcgtgc 300
cccacccatc caagccaaac cacaccctgg tccttctgga caccgaaggt ctgggcgatg 360
tggaaaaggg tgaccctaag aatgactcct ggatctttgc cctggctgtg ctcctgtgca 420
gcacctttgt ctacaacagc atgagcacca tcaaccacca gccctggag cagctgcatt 480
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atgtgacgga c   491

<210> 91
<211> 488
<212> DNA
<213> Homo sapiens

<400> 91
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ctgcttttaa ctctggtaaa gtggatattg ttgccatcaa tgaccccttc attgacctca 180
actacatggt ttacatgttc caatatgatt ccacccatgg caaattccat ggcaccgtcg 240
aggctgagaa cgggaagctt gtcatcaatg aaatcccat caccatcttc aggagcgag 300
atccctccaa aatcaagtgg ggcgatgctg cgctgagta cgtcgtggag tccactggcg 360
tcttcaccac catggagaag gctgggggctc atttgcaggg gggagccaaa agggtcatca 420
tctctgcccc tctgctgatg ccccatgttc gtcatgggtg tgaaccatga gaagtatgac 480
acagcctc   488

<210> 92
<211> 384
<212> DNA
<213> Homo sapiens

<400> 92
gacagtcagc cgcatcttct tttgcgtcgc cagccgagcc acatcgctca gacaccatgg 60
ggaaggtgaa ggtcggagtc aacggatttg tcgtattgg gcgcctggtc accagggctg 120
cttttaactc tggtaaagtg gatattgttg ccatcaatga ccccttcatt gacctcaact 180
acatggttta catgttccaa tatgattcca cccatggcaa attccatggc accgtcgagg 240
ctgagaacgg gaagcttgtc atcaatggaa atcccatcac catcttccag gagcgagatc 300
cctccaaaat caagtggggc gatactggcg ctgagtacgt cgtggagtcc actggcgtct 360
tcaccaccat ggagaaggct gggg   384

<210> 93
<211> 162
<212> PRT
<213> Homo sapiens

<400> 93
Lys Gly Lys Leu Asp Asp Tyr Gln Glu Arg Met Asn Lys Gly Glu Arg
1   5   10   15

Leu Asn Gln Asp Gln Leu Asp Ala Val Ser Lys Tyr Gln Glu Val Thr
                20   25   30

Asn Asn Leu Glu Phe Ala Lys Glu Leu Gln Arg Ser Phe Met Ala Leu
            35   40   45

Ser Gln Asp Ile Gln Lys Thr Ile Lys Lys Thr Ala Arg Arg Glu Gln
        50   55   60

Leu Met Arg Glu Glu Ala Glu Gln Lys Arg Leu Lys Thr Val Leu Glu
65   70   75   80

Leu Gln Tyr Val Leu Asp Lys Leu Gly Asp Asp Glu Val Arg Thr Asp
                85   90   95

```
Leu Lys Gln Gly Leu Asn Gly Val Pro Ile Leu Ser Glu Glu Glu Leu
          100               105               110

Ser Leu Leu Asp Glu Phe Tyr Lys Leu Val Asp Pro Glu Arg Asp Met
          115               120               125

Ser Leu Arg Leu Asn Glu Gln Tyr Glu His Ala Ser Ile His Leu Trp
      130                   135                 140

Asp Leu Leu Glu Gly Lys Glu Lys Pro Val Cys Gly Thr Thr Tyr Lys
145                   150               155                 160

Val Leu
```

<210> 94
<211> 100
<212> PRT
<213> Homo sapiens

<400> 94
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Asp Leu Glu Glu Ala Thr Leu Gln His Glu Ala Thr Ala Ala Thr Leu
  1               5               10                  15

Arg Lys Lys His Ala Asp Ser Val Ala Glu Leu Gly Glu Gln Ile Asp
            20                  25                  30

Asn Leu Gln Arg Val Lys Gln Lys Leu Glu Lys Glu Lys Ser Glu Met
            35                  40                  45

Lys Met Glu Ile Asp Asp Leu Ala Cys Asn Met Glu Val Ile Ser Lys
        50                  55                  60

Ser Lys Gly Asn Leu Glu Lys Met Cys Arg Thr Leu Glu Asp Gln Val
65                  70                  75                  80

Ser Glu Leu Lys Thr Gln Glu Glu Glu Gln Gln Arg Leu Ile Asn Glu
                85                  90                  95

Leu Thr Ala Gln
            100
```

<210> 95
<211> 99
<212> PRT
<213> Homo sapiens

<400> 95
```
Lys Ile Leu Pro Leu Asn Gly Asn Leu Gln Ala Val Glu Leu Gly Glu
  1               5               10                  15

Lys Arg Thr Ser Ser Leu Arg Ile Lys Met Phe Arg Ala Thr Arg Val
            20                  25                  30
```

```
Thr Ser Thr Ser Arg Phe Leu Asn Pro Tyr Val Val Cys Phe Leu Val
        35              40              45

Leu Pro Gly Val Val Ile Leu Ala Val Pro Ile Ala Leu Leu Val Tyr
    50              55              60

Phe Leu Ala Phe Asp Gln Lys Ser Tyr Phe Tyr Trp Ser Asn Phe Pro
65              70              75              80

Leu Pro Asn Val Glu Tyr Asn Ser Pro Phe Asn Ser Pro Ala Ser Pro
            85              90              95

Gly Ile Pro
```


<210> 96
<211> 257
<212> PRT
<213> Homo sapiens

<400> 96
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Val Gln Glu Thr Ile His Glu His Asn Lys Leu Ala Ala Asn Ser Asp
 1           5              10              15

His Leu Met Gln Ile Gln Lys Cys Glu Leu Val Leu Ile His Thr Tyr
            20              25              30

Pro Val Gly Glu Asp Ser Leu Val Ser Asp Arg Ser Lys Lys Glu Leu
        35              40              45

Ser Pro Val Leu Thr Ser Glu Val His Ser Val Arg Ala Gly Arg His
    50              55              60

Leu Ala Thr Lys Leu Asn Ile Leu Val Gln Gln His Phe Asp Leu Ala
65              70              75              80

Ser Thr Thr Ile Thr Asn Ile Pro Met Lys Glu Glu Gln His Ala Asn
            85              90              95

Thr Ser Ala Asn Tyr Asp Val Glu Leu Leu His His Lys Asp Ala His
            100             105             110

Val Asp Phe Leu Lys Ser Gly Asp Ser His Leu Gly Gly Gly Ser Arg
        115             120             125

Glu Gly Ser Phe Lys Glu Thr Ile Thr Leu Lys Trp Cys Thr Pro Arg
    130             135             140

Thr Asn Asn Ile Glu Leu His Tyr Cys Thr Gly Ala Tyr Arg Ile Ser
145             150             155             160

Pro Val Asp Val Asn Ser Arg Pro Ser Ser Cys Leu Thr Asn Phe Leu
            165             170             175
```

Leu Asn Gly Arg Ser Val Leu Leu Glu Gln Pro Arg Lys Ser Gly Ser
        180   185   190

Lys Val Ile Ser His Met Leu Ser Ser His Gly Gly Glu Ile Phe Leu
        195   200   205

His Val Leu Ser Ser Ser Arg Ser Ile Leu Glu Asp Pro Pro Ser Ile
     210   215   220

Ser Glu Gly Cys Gly Gly Arg Val Thr Asp Tyr Arg Ile Thr Asp Phe
225   230   235   240

Gly Glu Phe Met Arg Gly Lys Gln Ile Asn Ser Phe Ser Thr Pro Gln
            245   250   255

Ile

<210> 97
<211> 128
<212> PRT
<213> Homo sapiens

<400> 97
Ser Leu Pro Gln Phe Ala Val His Pro Glu Arg Ser Gly Leu Ala Asp
 1   5   10   15

Ser Gly Asp Gly Gly Asn Met Ser Val Ala Phe Ala Ala Pro Arg Gln
            20   25   30

Arg Gly Lys Gly Glu Ile Thr Pro Ala Ala Ile Gln Lys Met Leu Asp
        35   40   45

Asp Asn Asn His Leu Ile Gln Cys Ile Met Asp Ser Gln Asn Lys Gly
     50   55   60

Lys Thr Ser Glu Cys Ser Gln Tyr Gln Gln Met Leu His Thr Asn Leu
65   70   75   80

Val Tyr Leu Ala Thr Ile Ala Asp Ser Asn Gln Asn Met Gln Ser Leu
            85   90   95

Leu Pro Ala Pro Pro Thr Gln Asn Met Pro Met Gly Pro Gly Gly Met
            100   105   110

Asn Gln Ser Gly Pro Pro Pro Pro Arg Ser His Asn Met Pro Ser
        115   120   125

<210> 98
<211> 159
<212> PRT
<213> Homo sapiens

<400> 98
Phe Leu Asp Leu Arg Cys Tyr Arg Ala Gly Ser Ser Arg Leu Ala Val
 1   5   10   15

Ala Met Glu Ser Gly Pro Lys Met Leu Ala Pro Val Cys Leu Val Glu
            20   25   30

Asn Asn Asn Glu Gln Leu Leu Val Asn Gln Gln Ala Ile Gln Ile Leu
        35   40   45

Glu Lys Ile Ser Gln Pro Val Val Val Val Ala Ile Val Gly Leu Tyr
    50   55   60

Arg Thr Gly Lys Ser Tyr Leu Met Asn His Leu Ala Gly Gln Asn His
65   70   75   80

Gly Phe Pro Leu Gly Ser Thr Val Gln Ser Glu Thr Lys Gly Ile Trp
            85   90   95

Met Trp Cys Val Pro His Pro Ser Lys Pro Asn His Thr Leu Val Leu
            100   105   110

Leu Asp Thr Glu Gly Leu Gly Asp Val Glu Lys Gly Asp Pro Lys Asn
        115   120   125

Asp Ser Trp Ile Phe Ala Leu Ala Val Leu Leu Cys Ser Thr Phe Val
    130   135   140

Tyr Asn Ser Met Ser Thr Ile Asn His Gln Ala Leu Glu Gln Leu
145   150   155

<210> 99
<211> 147
<212> PRT
<213> Homo sapiens

<400> 99
Met Glu Ser Gly Pro Lys Met Leu Ala Pro Val Cys Leu Val Glu Asn
 1   5   10   15

Asn Asn Glu Gln Leu Leu Val Asn Gln Gln Ala Ile Gln Ile Leu Glu
            20   25   30

Lys Ile Ser Gln Pro Val Val Val Val Ala Ile Val Gly Leu Tyr Arg
        35   40   45

Thr Gly Lys Ser Tyr Leu Met Asn His Leu Ala Gly Gln Asn His Gly
    50   55   60

Phe Pro Leu Gly Ser Thr Val Gln Ser Glu Thr Lys Gly Ile Trp Met
65   70   75   80

Trp Cys Val Pro His Pro Ser Lys Pro Asn His Thr Leu Val Leu Leu

```
                        85                      90                      95
    Asp Thr Glu Gly Leu Gly Asp Val Glu Lys Gly Asp Pro Lys Asn Asp
                    100                     105                     110

    Ser Trp Ile Phe Ala Leu Ala Val Leu Leu Cys Ser Thr Phe Val Tyr
                    115                     120                     125

    Asn Ser Met Ser Thr Ile Asn His Gln Ala Leu Glu Gln Leu His Tyr
            130                     135                     140

    Val Thr Asp
    145
```

<210> 100
<211> 124
<212> PRT
<213> Homo sapiens

<400> 100

```
    Met Gly Lys Val Lys Val Gly Val Asn Gly Phe Gly Arg Ile Gly Arg
    1               5                   10                      15

    Leu Val Thr Arg Ala Ala Phe Asn Ser Gly Lys Val Asp Ile Val Ala
                    20                  25                      30

    Ile Asn Asp Pro Phe Ile Asp Leu Asn Tyr Met Val Tyr Met Phe Gln
            35                      40                      45

    Tyr Asp Ser Thr His Gly Lys Phe His Gly Thr Val Glu Ala Glu Asn
            50                      55                      60

    Gly Lys Leu Val Ile Asn Gly Asn Pro Ile Thr Ile Phe Gln Glu Arg
    65                  70                  75                      80

    Asp Pro Ser Lys Ile Lys Trp Gly Asp Ala Gly Ala Glu Tyr Val Val
                    85                  90                      95

    Glu Ser Thr Gly Val Phe Thr Thr Met Glu Lys Ala Gly Ala His Leu
                    100                 105                     110

    Gln Gly Gly Ala Lys Arg Val Ile Ile Ser Ala Pro
            115                     120
```

<210> 101
<211> 127
<212> PRT
<213> Homo sapiens

<400> 101

```
    Gln Ser Ala Ala Ser Ser Phe Ala Ser Pro Ala Glu Pro His Arg Ser
    1               5                   10                      15
```

```
Asp Thr Met Gly Lys Val Lys Val Gly Val Asn Gly Phe Gly Arg Ile
        20                  25                  30

Gly Arg Leu Val Thr Arg Ala Ala Phe Asn Ser Gly Lys Val Asp Ile
        35                  40                  45

Val Ala Ile Asn Asp Pro Phe Ile Asp Leu Asn Tyr Met Val Tyr Met
        50                  55                  60

Phe Gln Tyr Asp Ser Thr His Gly Lys Phe His Gly Thr Val Glu Ala
    65              70                  75                  80

Glu Asn Gly Lys Leu Val Ile Asn Gly Asn Pro Ile Thr Ile Phe Gln
            85                  90                  95

Glu Arg Asp Pro Ser Lys Ile Lys Trp Gly Asp Thr Gly Ala Glu Tyr
            100                 105                 110

Val Val Glu Ser Thr Gly Val Phe Thr Thr Met Glu Lys Ala Gly
        115                 120                 125
```

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<210> 102
<211> 1225
<212> DNA
<213> Homo sapiens

<400> 102
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gcggagacgg cagccgtgac ggtggcagcg gcggcgcggg acctgggcct gggggaatga 120
ggcggccgcg gcgggccagc ggcggagccg tgtagcggag aagctccccc tccctgcttc 180
ccttggccga gccggggggcg cgcgcgcacg cggccgtcca gagcgggctc cccacccctc 240
gactcctgcg acccgcaccg cacccccacc cgggcccgga ggatgatgaa gctcaagtcg 300
aaccagaccc gcacctacga cggcgacggc tacaagaagc gggccgcatg cctgtgtttc 360
cgcagcgaga gcgaggagga ggtgctactc gtgagcagta gtcgccatcc agacagatgg 420
attgtccctg gaggaggcat ggagcccgag gaggagccaa gtgtggcagc agttcgtgaa 480
gtctgtgagg aggctggagt aaaagggaca ttgggaagat tagttggaat ttttgagaac 540
caggagagga agcacaggac gtatgtctat gtgctcattg tcactgaagt gctggaagac 600
tgggaagatt cagttaacat tggaaggaag agggaatggt ttaaaataga agacgccata 660
aaagtgctgc agtatcacaa acccgtgcag gcatcatatt ttgaaacatt gaggcaaggc 720
tactcagcca acaatggcac cccagtcgtg gccaccacat actcggtttc tgctcagagc 780
tcgatgtcag gcatcagatg actgaagact tcctgtaaga gaaatggaaa ttggaaacta 840
gactgaagtg caaatcttcc ctctcaccct ggctctttcc acttctcaca ggcctcctct 900
ttcaaataag gcatggtggg cagcaaagaa agggtgtatt gataatgttg ctgtttggtg 960
ttaagtgatg gggcttttttc ttctgttttt attgagggtg ggggttgggt gtgtaatttg 1020
taagtacttt tgtgcatgat ctgtccctcc ctcttcccac ccctgcagtc ctctgaagag 1080
aggccaacag ccttcccctg ccttggattc tgaagtgttc ctgtttgtct tatcctggcc 1140
ctggccagac gtttttctttg attttttaatt ttttttttttt attaaaagat accagtatga 1200
gaaaaaaaaa aaaaaaaaac tcgag 1225
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<210> 103
<211> 741
<212> DNA
<213> Homo sapiens
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<400> 103
agaaacctca atcggattca gcaaaggaat ggtgttatta tcactacata ccaaatgtta 60
atcaataact ggcagcaact ttcaagcttt aggggccaag agtttgtgtg ggactatgtc 120
atcctcgatg aagcacataa aataaaaacc tcatctacta agtcagcaat atgtgctcgt 180
gctattcctg caagtaatcg cctcctcctc acaggaaccc caatccagaa taatttacaa 240
gaactatggt ccctatttga ttttgcttgt caagggtccc tgctgggaac attaaaaact 300
tttaagatgg agtatgaaaa tcctattact agagcaagag agaaggatgc taccccagga 360
gaaaaagcct tgggatttaa aatatctgaa aacttaatgg caatcataaa accctatttt 420
ctcaggagga ctaaagaaga cgtacagaag aaaaagtcaa gcaacccaga ggccagactt 480
aatgaaaaga atccagatgt tgatgccatt tgtgaaatgc cttccctttc caggagaaat 540
gatttaatta tttggatacg acttgtgcct ttacaagaag aaatatacag gaaatttgtg 600
tctttagatc atatcaagga gttgctaatg gagacgcgct caccttttgc tgagctaggt 660
gtcttaaaga agctgtgtga tcatcctagg ctgctgtctg cacgggcttg ttgtttgcta 720
aatcttggga cattctctgc t   741

<210> 104
<211> 321
<212> DNA
<213> Homo sapiens

<400> 104
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cctcagatgg aactgccact ccaaggctgt aacattacgt acatcccgaa agacagcaaa 120
aagaagaagc acgagctgaa gattactcag cagggcacgg acccgcttgt tctcgccgtc 180
cagagcaagg aacaggccga gcagtggctg aaggtgatca agaagcctta cagtggttgt 240
agtggccccg tggattcaga gtgtcctcct ccaccaagct ccccggtgca caaggcagaa 300
ctggagaaga aactgtcttc a   321

<210> 105
<211> 389
<212> DNA
<213> Homo sapiens

<400> 105
cagcactggc cacactataa aattcaggtt cagaaaaaca gggtaagtca cagacagcaa 60
cgcttccagc atttattttc tttgcaccca tgggcaattc gagaaaattt acctttagaa 120
cgaactctgt taaaggtaca gacagtacaa tactttttat tcagaaggtt tctgcataaa 180
ggtgatagtc ttttgactta atatattatt gtctcctgcc ttgtgtttct ggaatgaatg 240
aaggtcatta tttagaagat aatctgggtt gtatttgtgt cgtcagattg aattttcatt 300
gcacatgcta cttaatgtct ttaccaaata ataacaaagg gaaagaaaac caaatataga 360
tgtataataa ggaaaagctg gcctataga   389

<210> 106
<211> 446
<212> DNA
<213> Homo sapiens

<400> 106
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acaagtatca ctccattgtt cagagagtaa tgtattagtt ctgcccaatt cattcttcac 120
ttttatttct tccatttcat tagcatttat atcagctcaa gaagttaagg ttagaaaatt 180
ttccacttca aatttttcagt acagaaatgt gctgtgatgt ttgacaagac tatttcatag 240
taagtgagtt aatgtttatt ggcctctgct ctcctctgtg tcagacctag gaagcctgag 300
gattacttag ttgttctgtc tctgggtcca caggcagaat ttggcccatc caaagactgg 360
ccaagtgcca aaaaaaggcc tgattaggcc ctgaaattca gtgaaattct gcctgaagaa 420

acctcttatt gaatttgaaa accata   446

<210> 107
<211> 467
<212> DNA
<213> Homo sapiens

<400> 107
ccgccgctgc cgtcgccttc ctgggattgg agtctcgagc tttcttcgtt cgttcgccgg 60
cgggttcgcg cccttctcgc gcctcggggc tgcgaggctg gggaaggggt tggaggggggc 120
tgttgatcgc cgcgtttaag ttgcgctcgg ggcggccatg tcggccggcg aggtcgagcg 180
cctagtgtcg gagctgagcg gcgggaccgg aggggatgag gaggaagagt ggctctatgg 240
cgatgaagat gaagttgaaa ggccagaaga agaaaatgcc agtgctaatc ctccatctgg 300
aattgaagat gaaactgctg aaaatggtgt accaaaaccg aaagtgactg agaccgaaga 360
tgatagtgat agtgacagcg atgatgatga agatgatgtg catgtcacta taggagacat 420
taaaacggga gcaccacagt atgggagtta tggtacagca cctgtaa   467

<210> 108
<211> 491
<212> DNA
<213> Homo sapiens

<400> 108
gaaagataca acttccccaa cccaaacccg tttgtggagg acgacatgga taagaatgaa 60
atcgcctctg ttgcgtaccg ttaccgcagg tggaagcttg gagatgatat tgaccttatt 120
gtccgttgtg agcacgatgg cgtcatgact ggagccaacg gggaagtgtc cttcatcaac 180
atcaagacac tcaatgagtg ggattccagg cactgtaatg gcgttgactg gcgtcagaag 240
ctggactctc agcgaggggc tgtcattgcc acggagctga agaacaacag ctacaagttg 300
gcccggtgga cctgctgtgc tttgctggct ggatctgagt acctcaagct tggttatgtg 360
tctcggtacc acgtgaaaga ctcctcacgc cacgtcatcc taggcaccca gcagttcaag 420
cctaatgagt ttgccagcca gatcaacctg agcgtggaga atgcctgagg cattttacgc 480
tgcgtcattg a   491

<210> 109
<211> 489
<212> DNA
<213> Homo sapiens

<400> 109
ctcagatagt actgaaccct ttatcaacta tgttttttca gtctgacaac caaggcggct 60
actaagtgac taaggggcag gtagtataca gtgtggataa gcaggacaaa ggggtgattc 120
acatcccagg caggacagag caggagatca tgagatttca tcactcagga tggcttgtga 180
tttatttat tttattcttt ttttttttttg agatggagtc tcactcttgc ccaggctgga 240
gtgcagtggt gcgatcttgg ctcactgcaa cctctgcctc ctgggttcaa gcagttctcc 300
tgcctcagcc tcccaagtag ctgggattac aggcgtccgc caccatgccc agccaatttt 360
tgtacttta gtagagatgg ggtttcacca tgttggccag gctggtctcg aactcctgac 420
ctcaggtgat ccactcgcct cggcctccca aagtgctggg attataggca tgcgccacca 480
tgcccgggc   489

<210> 110
<211> 391
<212> DNA
<213> Homo sapiens

<400> 110

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gcggagtccg ctggctgacc cgagcgctgg tctccgccgg gaaccctggg gcatggagag 60
gtctgagtac ctcggccgcg gcgcacgctg catcgcggag ccaggctgcc gctgtcccag 120
tggagttcca ggagcaccac ctgagtgagg tgcagaatat ggcatctgag gagaagctgg 180
agcaggtgct gagttccatg aaggagaaca aagtggccat cattggaaag attcataccc 240
cgatggagta taaggggggag ctagcctcct atgatatgcg gctgaggcgt aagttggact 300
tatttgccaa cgtaatccat gtgaagtcac ttcctgggta tatgactcgg cacaacaatc 360
tagacctggt gatcattcga gagcagacag a                             391
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<210> 111
<211> 172
<212> PRT
<213> Homo sapiens

<400> 111
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Met Met Lys Leu Lys Ser Asn Gln Thr Arg Thr Tyr Asp Gly Asp Gly
 1               5                   10                  15

Tyr Lys Lys Arg Ala Ala Cys Leu Cys Phe Arg Ser Glu Ser Glu Glu
                20              25              30

Glu Val Leu Leu Val Ser Ser Ser Arg His Pro Asp Arg Trp Ile Val
         35                  40                  45

Pro Gly Gly Gly Met Glu Pro Glu Glu Glu Pro Ser Val Ala Ala Val
     50                  55                  60

Arg Glu Val Cys Glu Glu Ala Gly Val Lys Gly Thr Leu Gly Arg Leu
 65                  70                  75                  80

Val Gly Ile Phe Glu Asn Gln Glu Arg Lys His Arg Thr Tyr Val Tyr
                 85                  90                  95

Val Leu Ile Val Thr Glu Val Leu Glu Asp Trp Glu Asp Ser Val Asn
                100                 105                 110

Ile Gly Arg Lys Arg Glu Trp Phe Lys Ile Glu Asp Ala Ile Lys Val
            115                 120                 125

Leu Gln Tyr His Lys Pro Val Gln Ala Ser Tyr Phe Glu Thr Leu Arg
        130                 135                 140

Gln Gly Tyr Ser Ala Asn Asn Gly Thr Pro Val Val Ala Thr Thr Tyr
    145                 150                 155                 160

Ser Val Ser Ala Gln Ser Ser Met Ser Gly Ile Arg
                165                 170
```

<210> 112
<211> 247
<212> PRT
<213> Homo sapiens

<400> 112
Arg Asn Leu Asn Arg Ile Gln Gln Arg Asn Gly Val Ile Ile Thr Thr

```
         1               5                    10                   15 .

      Tyr Gln Met Leu Ile Asn Asn Trp Gln Gln Leu Ser Ser Phe Arg Gly
                    20                  25                  30

      Gln Glu Phe Val Trp Asp Tyr Val Ile Leu Asp Glu Ala His Lys Ile
                  35                40                  45

      Lys Thr Ser Ser Thr Lys Ser Ala Ile Cys Ala Arg Ala Ile Pro Ala
             50                    55                  60

      Ser Asn Arg Leu Leu Leu Thr Gly Thr Pro Ile Gln Asn Asn Leu Gln
      65              70                  75                      80

      Glu Leu Trp Ser Leu Phe Asp Phe Ala Cys Gln Gly Ser Leu Leu Gly
                      85                  90                      95

      Thr Leu Lys Thr Phe Lys Met Glu Tyr Glu Asn Pro Ile Thr Arg Ala
                  100                 105                 110

      Arg Glu Lys Asp Ala Thr Pro Gly Glu Lys Ala Leu Gly Phe Lys Ile
               115                 120                 125

      Ser Glu Asn Leu Met Ala Ile Ile Lys Pro Tyr Phe Leu Arg Arg Thr
           130                 135                 140

      Lys Glu Asp Val Gln Lys Lys Lys Ser Ser Asn Pro Glu Ala Arg Leu
      145                 150                 155                 160

      Asn Glu Lys Asn Pro Asp Val Asp Ala Ile Cys Glu Met Pro Ser Leu
                     165                 170                 175

      Ser Arg Arg Asn Asp Leu Ile Ile Trp Ile Arg Leu Val Pro Leu Gln
                 180                 185                 190

      Glu Glu Ile Tyr Arg Lys Phe Val Ser Leu Asp His Ile Lys Glu Leu
               195                 200                 205

      Leu Met Glu Thr Arg Ser Pro Leu Ala Glu Leu Gly Val Leu Lys Lys
      210                 215                 220

      Leu Cys Asp His Pro Arg Leu Leu Ser Ala Arg Ala Cys Cys Leu Leu
      225                 230                 235                 240

      Asn Leu Gly Thr Phe Ser Ala
                      245
```

<210> 113
<211> 107
<212> PRT
<213> Homo sapiens

<400> 113
Leu Leu Cys Val Ile Lys Asp Thr Lys Leu Leu Cys Tyr Lys Ser Ser

```
        1               5                   10                  15
      Lys Asp Gln Gln Pro Gln Met Glu Leu Pro Leu Gln Gly Cys Asn Ile
                  20                  25                  30

      Thr Tyr Ile Pro Lys Asp Ser Lys Lys Lys His Glu Leu Lys Ile
                  35                  40                  45

      Thr Gln Gln Gly Thr Asp Pro Leu Val Leu Ala Val Gln Ser Lys Glu
            50                  55                  60

      Gln Ala Glu Gln Trp Leu Lys Val Ile Lys Glu Ala Tyr Ser Gly Cys
      65                  70                  75                  80

      Ser Gly Pro Val Asp Ser Glu Cys Pro Pro Pro Ser Ser Pro Val
                  85                  90                  95

      His Lys Ala Glu Leu Glu Lys Lys Leu Ser Ser
                  100                 105
```

```
<210> 114
<211> 155
<212> PRT
<213> Homo sapiens

<400> 114
      Glu Arg Tyr Asn Phe Pro Asn Pro Asn Pro Phe Val Glu Asp Asp Met
       1               5                   10                  15

      Asp Lys Asn Glu Ile Ala Ser Val Ala Tyr Arg Tyr Arg Arg Trp Lys
                  20                  25                  30

      Leu Gly Asp Asp Ile Asp Leu Ile Val Arg Cys Glu His Asp Gly Val
                  35                  40                  45

      Met Thr Gly Ala Asn Gly Glu Val Ser Phe Ile Asn Ile Lys Thr Leu
            50                  55                  60

      Asn Glu Trp Asp Ser Arg His Cys Asn Gly Val Asp Trp Arg Gln Lys
      65                  70                  75                  80

      Leu Asp Ser Gln Arg Gly Ala Val Ile Ala Thr Glu Leu Lys Asn Asn
                  85                  90                  95

      Ser Tyr Lys Leu Ala Arg Trp Thr Cys Cys Ala Leu Leu Ala Gly Ser
                  100                 105                 110

      Glu Tyr Leu Lys Leu Gly Tyr Val Ser Arg Tyr His Val Lys Asp Ser
                  115                 120                 125

      Ser Arg His Val Ile Leu Gly Thr Gln Gln Phe Lys Pro Asn Glu Phe
            130                 135                 140

      Ala Ser Gln Ile Asn Leu Ser Val Glu Asn Ala
```

145   150   155

<210> 115
<211> 129
<212> PRT
<213> Homo sapiens

<400> 115
Gly Val Arg Trp Leu Thr Arg Ala Leu Val Ser Ala Gly Asn Pro Gly
 1   5   10   15

Ala Trp Arg Gly Leu Ser Thr Ser Ala Ala Ala His Ala Ala Ser Arg
            20   25   30

Ser Gln Ala Ala Ala Val Pro Val Glu Phe Gln Glu His His Leu Ser
        35   40   45

Glu Val Gln Asn Met Ala Ser Glu Glu Lys Leu Glu Gln Val Leu Ser
    50   55   60

Ser Met Lys Glu Asn Lys Val Ala Ile Ile Gly Lys Ile His Thr Pro
65   70   75   80

Met Glu Tyr Lys Gly Glu Leu Ala Ser Tyr Asp Met Arg Leu Arg Arg
                85   90   95

Lys Leu Asp Leu Phe Ala Asn Val Ile His Val Lys Ser Leu Pro Gly
            100   105   110

Tyr Met Thr Arg His Asn Asn Leu Asp Leu Val Ile Ile Arg Glu Gln
            115   120   125

Thr

<210> 116
<211> 550
<212> DNA
<213> Homo sapiens

<400> 116
gaattcggca ccagcctcag agccccccag cccggctacc accccctgcg gaaaggtacc 60
catctgcatt cctgcccgtc gggacctggt ggacagtcca gcctccttgg cctctagcct 120
tggctcaccg ctgcctagag ccaaggagct catcctgaat gaccttcccg ccagcactcc 180
tgcctccaaa tcctgtgact cctcccgcc ccaggacgct tccacccca ggcccagctc 240
ggccagtcac ctctgccagc ttgctgccaa gccagcacct tccacggaca gcgtcgccct 300
gaggagcccc ctgactctgt ccagtccctt caccacgtcc ttcagcctgg ctcccacag 360
cactctcaac ggagacctct ccgtgcccag ctcctacgtc agcctccacc tgtcccccca 420
ggtcagcagc tctgtggtgt acggacgctc ccccgtgatg gcatttgagt ctcatcccca 480
tctccgaggg tcatccgtct cttcctccct acccagcatc cctggggggaa agccggccta 540
ctccttccac   550

<210> 117
<211> 154

<212> DNA
<213> Homo sapiens

<400> 117
ttctgaggga aagccgagtg gagtgggcga cccggcggcg gtgacaatga gttttcttgg 60
aggctttttt ggtcccattt gtgagattga tgttgccctt aatgatgggg aaaccaggaa 120
aatggcagaa atgaaaactg aggatggcaa agta   154

<210> 118
<211> 449
<212> DNA
<213> Homo sapiens

<400> 118
gaattcggca ccagggcccg cagcccgagt gtcgccgcca tggcttcgcc gcagctctgc 60
cgcgcgctgg tgtcggcgca atgggtggcg gaggcgctgc gggccccgcg cgctgggcag 120
cctctgcagc tgctggacgc ctcctggtac ctgccgaagc tggggcgcga cgcgcgacgc 180
gagttcgagg agcgccacat cccgggcgcc gctttcttcg acatcgacca gtgcagcgac 240
cgcacctcgc cctacgacca catgctgccc ggggccgagc atttcgcgga gtacgcaggc 300
cgcctgggcg tgggcgcggc cacccacgtc gtgatctacg acgccagcga ccagggcctc 360
tactccgccc cgcgcgtctg gtggatgttc cgcgccttcg gccaccacgc cgtgtcactg 420
cttgatggcg gcctccgcca ctggctgcg   449

<210> 119
<211> 642
<212> DNA
<213> Homo sapiens

<400> 119
gaattcggca cgagcagtaa cccgaccgcc gctggtcttc gctggacacc atgaatcaca 60
ctgtccaaac cttcttctct cctgtcaaca gtggccagcc ccccaactat gagatgctca 120
aggaggagca cgaggtggct gtgctggggg cgccccacaa ccctgctccc ccgacgtcca 180
ccgtgatcca catccgcagc gagacctccg tgcccgacca tgtcgtctgg tccctgttca 240
acaccctctt catgaacccc tgctgcctgg gcttcatagc attcgcctac tccgtgaagt 300
ctagggacag gaagatggtt ggcgacgtga ccggggccca ggcctatgcc tccaccgcca 360
agtgcctgaa catctgggcc ctgattctgg gcatcctcat gaccattctg ctcatcgtca 420
tcccagtgct gatcttccag gcctatggat agatcaggag gcatcactga ggccaggagc 480
tctgcccatg acctgtatcc cacgtactcc aacttccatt cctcgccctg cccccggagc 540
cgagtcctgt atcagcccct tatcctcaca cgcttttcta caatggcatt caataaagtg 600
cacgtgtttc tggtgaaaaa aaaaaaaaaa aaaaaactcg ag   642

<210> 120
<211> 603
<212> DNA
<213> Homo sapiens

<400> 120
gaattcggca cgagccacaa cagccactac gactgcatcc actggatcca cggccacccc 60
gtcctccacc ccgggaacag ctcccccctcc caaagtgctg accagcccgg ccaccacacc 120
catgtccacc atgtccacaa tccacacctc ctctactcca gagaccaccc acacctccac 180
agtgctgacc accacagcca ccatgacaag ggccaccaat tccacggcca caccctcctc 240
cactctgggg acgacccgga tcctcactga gctgaccaca acagccacta caactgcagc 300
cactggatcc acggccaccc tgtcctccac cccagggacc acctggatcc tcacagagcc 360
gagcactata gccaccgtga tggtgcccac cggttccacg gccaccgcct cctccactct 420
gggaacagct cacaccccca aagtggtgac caccatggcc actatgccca cagccactgc 480

```
ctccacggtt cccagctcgt ccaccgtggg gaccacccgc acccctgcag tgctccccag 540
cagcctgcca accttcagcg tgtccactgt gtcctcctca gtcctcacca ccctgagacc 600
cac                                                                603
```

```
<210> 121
<211> 178
<212> PRT
<213> Homo sapiens
```

```
<400> 121
Ser Glu Pro Pro Ser Pro Ala Thr Thr Pro Cys Gly Lys Val Pro Ile
 1               5                  10                  15

Cys Ile Pro Ala Arg Arg Asp Leu Val Asp Ser Pro Ala Ser Leu Ala
            20                  25                  30

Ser Ser Leu Gly Ser Pro Leu Pro Arg Ala Lys Glu Leu Ile Leu Asn
        35                  40                  45

Asp Leu Pro Ala Ser Thr Pro Ala Ser Lys Ser Cys Asp Ser Ser Pro
    50                  55                  60

Pro Gln Asp Ala Ser Thr Pro Arg Pro Ser Ser Ala Ser His Leu Cys
65                  70                  75                  80

Gln Leu Ala Ala Lys Pro Ala Pro Ser Thr Asp Ser Val Ala Leu Arg
                85                  90                  95

Ser Pro Leu Thr Leu Ser Ser Pro Phe Thr Thr Ser Phe Ser Leu Gly
            100                 105                 110

Ser His Ser Thr Leu Asn Gly Asp Leu Ser Val Pro Ser Ser Tyr Val
            115                 120                 125

Ser Leu His Leu Ser Pro Gln Val Ser Ser Ser Val Val Tyr Gly Arg
    130                 135                 140

Ser Pro Val Met Ala Phe Glu Ser His Pro His Leu Arg Gly Ser Ser
145                 150                 155                 160

Val Ser Ser Ser Leu Pro Ser Ile Pro Gly Gly Lys Pro Ala Tyr Ser
            165                 170                 175

Phe His
```

```
<210> 122
<211> 36
<212> PRT
<213> Homo sapiens
```

```
<400> 122
Met Ser Phe Leu Gly Gly Phe Phe Gly Pro Ile Cys Glu Ile Asp Val
 1               5                  10                  15
```

```
Ala Leu Asn Asp Gly Glu Thr Arg Lys Met Ala Glu Met Lys Thr Glu
          20                25                30

Asp Gly Lys Val
          35


<210> 123
<211> 136
<212> PRT
<213> Homo sapiens

<400> 123
Met Ala Ser Pro Gln Leu Cys Arg Ala Leu Val Ser Ala Gln Trp Val
 1             5                10                15

Ala Glu Ala Leu Arg Ala Pro Arg Ala Gly Gln Pro Leu Gln Leu Leu
          20                25                30

Asp Ala Ser Trp Tyr Leu Pro Lys Leu Gly Arg Asp Ala Arg Arg Glu
          35                40                45

Phe Glu Glu Arg His Ile Pro Gly Ala Ala Phe Phe Asp Ile Asp Gln
     50                55                60

Cys Ser Asp Arg Thr Ser Pro Tyr Asp His Met Leu Pro Gly Ala Glu
 65                70                75                80

His Phe Ala Glu Tyr Ala Gly Arg Leu Gly Val Gly Ala Ala Thr His
               85                90                95

Val Val Ile Tyr Asp Ala Ser Asp Gln Gly Leu Tyr Ser Ala Pro Arg
          100               105               110

Val Trp Trp Met Phe Arg Ala Phe Gly His His Ala Val Ser Leu Leu
     115               120               125

Asp Gly Gly Leu Arg His Trp Leu
     130               135


<210> 124
<211> 133
<212> PRT
<213> Homo sapiens

<400> 124
Met Asn His Thr Val Gln Thr Phe Phe Ser Pro Val Asn Ser Gly Gln
 1             5                10                15

Pro Pro Asn Tyr Glu Met Leu Lys Glu Glu His Glu Val Ala Val Leu
          20                25                30

Gly Ala Pro His Asn Pro Ala Pro Pro Thr Ser Thr Val Ile His Ile
          35                40                45
```

Arg Ser Glu Thr Ser Val Pro Asp His Val Val Trp Ser Leu Phe Asn
    50   55   60

Thr Leu Phe Met Asn Pro Cys Cys Leu Gly Phe Ile Ala Phe Ala Tyr
    65   70   75   80

Ser Val Lys Ser Arg Asp Arg Lys Met Val Gly Asp Val Thr Gly Ala
                85   90   95

Gln Ala Tyr Ala Ser Thr Ala Lys Cys Leu Asn Ile Trp Ala Leu Ile
            100   105   110

Leu Gly Ile Leu Met Thr Ile Leu Leu Ile Val Ile Pro Val Leu Ile
            115   120   125

Phe Gln Ala Tyr Gly
    130

<210> 125
<211> 195
<212> PRT
<213> Homo sapiens

<400> 125
Thr Thr Ala Thr Thr Thr Ala Ser Thr Gly Ser Thr Ala Thr Pro Ser
1   5   10   15

Ser Thr Pro Gly Thr Ala Pro Pro Pro Lys Val Leu Thr Ser Pro Ala
            20   25   30

Thr Thr Pro Met Ser Thr Met Ser Thr Ile His Thr Ser Ser Thr Pro
            35   40   45

Glu Thr Thr His Thr Ser Thr Val Leu Thr Thr Thr Ala Thr Met Thr
    50   55   60

Arg Ala Thr Asn Ser Thr Ala Thr Pro Ser Ser Thr Leu Gly Thr Thr
65   70   75   80

Arg Ile Leu Thr Glu Leu Thr Thr Thr Ala Thr Thr Thr Ala Ala Thr
                85   90   95

Gly Ser Thr Ala Thr Leu Ser Ser Thr Pro Gly Thr Thr Trp Ile Leu
            100   105   110

Thr Glu Pro Ser Thr Ile Ala Thr Val Met Val Pro Thr Gly Ser Thr
            115   120   125

Ala Thr Ala Ser Ser Thr Leu Gly Thr Ala His Thr Pro Lys Val Val
    130   135   140

Thr Thr Met Ala Thr Met Pro Thr Ala Thr Ala Ser Thr Val Pro Ser
145   150   155   160

Ser Ser Thr Val Gly Thr Thr Arg Thr Pro Ala Val Leu Pro Ser Ser
              165   170   175

Leu Pro Thr Phe Ser Val Ser Thr Val Ser Ser Ser Val Leu Thr Thr
              180   185   190

Leu Arg Pro
       195

         <210> 126
         <211> 509
         <212> DNA
         <213> homo sapien

         <400> 126
    gaattcggca cgagccaagt accccctgag gaatctgcag cctgcatctg agtacaccgt   60
    atccctcgtg gccataaagg gcaaccaaga gagccccaaa gccactggag tctttaccac   120
    actgcagcct gggagctcta ttccacctta caacaccgag gtgactgaga ccaccattgt   180
    gatcacatgg acgcctgctc caagaattgg ttttaagctg ggtgtacgac caagccaggg   240
    aggagaggca ccacgagaag tgacttcaga ctcaggaagc atcgttgtgt ccggcttgac   300
    tccaggagta gaatacgtct acaccatcca agtcctgaga gatggacagg aaagagatgc   360
    gccaattgta aacaaagtgg tgacaccatt gtctccacca acaaacttgc atctggaggc   420
    aaaccctgac actggagtgc tcacagtctc ctggagagga gcaccacccc agacattact   480
    gggtatagaa ttaccacaac ccctacaaa   509

         <210> 127
         <211> 500
         <212> DNA
         <213> homo sapien

         <400> 127
    gaattcggca cgagccactg atgtccgggg agtcagccag gagcttgggg aagggaagcg   60
    cgcccccggg gccggtcccg gagggctcga tccgcatcta cagcatgagg ttctgcccgt   120
    ttgctgagag gacgcgtcta gtcctgaagg ccaagggaat caggcatgaa gtcatcaata   180
    tcaacctgaa aaataagcct gagtggttct ttaagaaaaa tccctttggt ctggtgccag   240
    ttctggaaaa cagtcagggt cagctgatct acgagtctgc catcacctgt gagtacctgg   300
    atgaagcata cccagggaag aagctgcttgc cggatgaccc ctatgagaaa gcttgccaga   360
    agatgatctt agagttgttt tctaaggtgc catccttggt aggaagcttt attagaagcc   420
    aaaataaaga agactatgct ggcctaaaag aagaatttcg taaagaattt accaagctag   480
    aggaggttct gactaataag   500

         <210> 128
         <211> 500
         <212> DNA
         <213> homo sapien

         <400> 128
    agctttcctc tgctgccgct cggtcacgct tgtgcccgaa ggaggaaaca gtgacagacc   60
    tggagactgc agttctctat ccttcacaca gctctttcac catgcctgga tcacttcctt   120
    tgaatgcaga agcttgctgg ccaaaagatg tgggaattgt tgcccttgag atctattttc   180
    cttctcaata tgttgatcaa gcagagttgg aaaaatatga tggtgtagat gctggaaagt   240
    ataccattgg cttgggccag gccaagatgg gcttctgcac agatagagaa gatattaact   300
    ctctttgcat gactgtggtt cagaatctta tggagagaaa taaccttttcc tatgattgca   360

```
ttgggcggct ggaagttgga acagagacaa tcatcgacaa atcaaagtct gtgaagacta        420
atttgatgca gctgtttgaa gagtctggga atacagatat agaaggaatc gacacaacta        480
atgcatgcta tggaggcaca                                                     500
```

        <210> 129
        <211> 497
        <212> DNA
        <213> homo sapien

        <400> 129
```
gaattcggca cgagcagagg tctccagagc cttctctctc ctgtgcaaaa tggcaactct         60
taaggaaaaa ctcattgcac cagttgcgga agaagaggca acagttccaa acaataagat        120
cactgtagtg ggtgttggac aagttggtat ggcgtgtgct atcagcattc tgggaaagtc        180
tctggctgat gaacttgctc ttgtggatgt tttggaagat aagcttaaag gagaaatgat        240
ggatctgcag catgggagct tatttcttca gacacctaaa attgtggcag ataaagatta        300
ttctgtgacc gccaattcta agattgtagt ggtaactgca ggagtccgtc agcaagaagg        360
ggagagtcgg ctcaatctgg tgcagagaaa tgttaatgtc ttcaaattca ttattcctca        420
gatcgtcaag tacagtcctg attgcatcat aattgtggtt tccaacccag tggacattct        480
tacgtatgtt acctgga                                                       497
```

        <210> 130
        <211> 383
        <212> DNA
        <213> homo sapien

        <400> 130
```
gaattcggca cgagggccgc ggctgccgac tgggtcccct gccgctgtcg ccaccatggc         60
tccgcaccgc cccgcgcccg cgctgctttg cgcgctgtcc ctggcgctgt gcgcgctgtc        120
gctgcccgtc cgcgcggcca ctgcgtcgcg ggggcgtcc caggcggggg cgccccaggg        180
gcgggtgccc gaggcgcggc ccaacagcat ggtggtggaa caccccgagt tcctcaaggc        240
agggaaggag cctggcctgc agatctggcg tgtggagaaa gttcgatctg gtggccgtg         300
cccaccaacc tttatggaga cttcttcacg ggcgacgcct acgtcatcct gaagacagtg        360
cagcttaaga acggaaaatc ttg                                                383
```

        <210> 131
        <211> 509
        <212> DNA
        <213> homo sapien

        <400> 131
```
gaattcggca cgagagtcag ccgcatcttc ttttgcgtcg ccagccgagc cacatcgctc         60
agacaccatg gggaaggtga aggtcggagt caacggattt ggtcgtattg ggcgcctggt        120
caccagggct gcttttaact ctggtaaagt ggatattgtt gccatcaatg acccccttcat       180
tgacctcaac tacatggttt acatgttcca atatgattcc acccatggca aattccatgg        240
caccgtcaag gctgagaacg ggaagcttgt catcaatgga aatcccatca ccatcttcca        300
ggagcgagat ccctccaaaa tcaagtgggg cgatgctggc gctgagtacg tcgtggagtc        360
cactggccgt cttcaccacc atggagaagg ctggggctca tttgcagggg ggagccaaaa        420
gggtcatcat ctctgccccc tctgctgacg cccccatgtt cgtcatgggt gtgaaccatg        480
agaagtatga caacagcctc aagatcatc                                          509
```

        <210> 132
        <211> 357
        <212> DNA
        <213> homo sapien

<400> 132
gaattcggca cgagtaagaa gaagccccta gaccacagct ccacaccatg gactggacct   60
ggaggatcct cttcttggtg gcagcagcaa caggtgccca ctcccaggtg caactggtgc   120
aatctgggtc tgagttgaag aagcctgggg cctcagtgaa ggtttcctgc aaggcttctg   180
gacacatctt cagtatctat ggtttgaatt gggtgcgaca ggcccctggt caaggccttg   240
agtggatggg atggatcaaa gtcgacactg cgaacccaac gtatgcccag ggcttcacag   300
gacgatttgt cttctccctg gacacctctg tcagcacggc atatctgcag atcagca   357

        <210> 133
        <211> 468
        <212> DNA
        <213> homo sapien

        <400> 133
gaattcggca cgaggcgccc cgaaccgtcc tcctgctgct ctcggcggcc ctggccctga   60
ccgagacctg ggccggctcc cactccatga ggtatttcga caccgccatg tcccggcccg   120
gccgcgggga gccccgcttc atctcagtgg gctacgtgga cgacacgcag ttcgtgaggt   180
tcgacagcga cgccgcgagt ccgagagagg agccgcgggc gccgtggata gagcaggagg   240
ggccggagta ttgggaccgg aacacacaga tcttcaagac caacacacag actgaccgag   300
agagcctgcg gaacctgcgc ggctactaca accagagcga ggccgggtct cacaccctcc   360
agagcatgta cggctgcgac gtggggccgg acgggcgcct cctccgcggg cataaccagt   420
acgcctacga cggcaaggat tacatcgccc tgaacgagga cctgcgct   468

        <210> 134
        <211> 214
        <212> DNA
        <213> homo sapien

        <400> 134
gaattcggca cgagctgcgt cctgctgagc tctgttctct ccagcacctc ccaacccact   60
agtgcctggt tctcttgctc caccaggaac aagccaccat gtctcgccag tcaagtgtgt   120
ccttccggag cggggggcagt cgtagcttca gcaccgcctc tgccatcacc ccgtctgtct   180
cccgcaccag cttcacctcc gtgtcccggt ccgg   214

        <210> 135
        <211> 355
        <212> DNA
        <213> homo sapien

        <400> 135
gaattcggca cgaggtgaac aggacccgtc gccatgggcc gtgtgatccg tggacagagg   60
aagggcgccg ggtctgtgtt ccgcgcgcac gtgaagcacc gtaaaggcgc tgcgcgcctg   120
cgcgccgtgg atttcgctga gcggcacggc tacatcaagg gcatcgtcaa ggacatcatc   180
cacgacccgg gccgcggcgc gcccctcgcc aaggtggtct tccgggatcc gtatcggttt   240
aagaagcgga cggagctgtt cattgccgcc gagggcattc acacgggcca gtttgtgtat   300
tgcggcaaga aggcccagct caacattggc aatgtgctcc ctgtgggcac catgc   355

        <210> 136
        <211> 242
        <212> DNA
        <213> homo sapien

        <400> 136
gaattcggca cgagccagct cctaaccgcg agtgatccgc cagcctccgc ctcccgaggt   60
gcccggattg cagacggagt ctccttcact cagtgctcaa tggtgcccag gctggagtgc   120

```
agtggtgtga tctcggctcg ctacaacatc cacctcccag cagcctgcct tggcctccca     180
aagtgccgag attgcagctc tctgcccggc cgccacccct gtctgggaag tgaggatgct     240
gt                                                                    242
```

```
        <210> 137
        <211> 424
        <212> DNA
        <213> homo sapien
```

```
        <400> 137
gaattcggca cgagcccaga tcccgaggtc cgacagcgcc cggcccagat ccccacgcct      60
gccaggagca agccgagagc cagccggccg gcgcactccg actccgagca gtctctgtcc     120
ttcgacccga gccccgcgcc ctttccggga cccctgcccc gcgggcagcg ctgccaacct     180
gccggccatg gagacccgt cccagcggcg cgccacccgc agcggggcgc aggccagctc      240
cactccgctg tcgcccaccc gcatcacccg gctgcaggag aaggaggacc tgcaggagct     300
caatgatcgc ttggcggtct acatcgaccg tgtgcgctcg ctggaaacgg agaacgcagg     360
gctgcgcctt cgcatcaccg agtctgaaga ggtggtcagc cgcgaggtgt ccggcatcaa     420
ggcc                                                                 424
```

```
        <210> 138
        <211> 448
        <212> DNA
        <213> homo sapien
```

```
        <400> 138
gaattcggca cgagcctgtg ttccaggagc cgaatcagaa atgtcatcct caggcacgcc      60
agacttacct gtcctactca ccgatttgaa gattcaatat actaagatct tcataaacaa     120
tgaatggcat gattcagtga gtggcaagaa atttcctgtc tttaatcctg caactgagga     180
ggagctctgc caggtagaag aaggagataa ggaggatgtt gacaaggcag tgaaggccgc     240
aagacaggct tttcagattg gatccccgtg gcgtactatg gatgcttccg agaggggggcg     300
actattatac aagttggctg atttaatcga aagagatcgt ctgctgctgg ccgacaatgg      360
agtcaatgaa tggtggaaaa ctctattcca atgcatatct gaatgattta gcaggctgca     420
tcaaaacatt gcgctactgt gcaggttg                                        448
```

```
        <210> 139
        <211> 510
        <212> DNA
        <213> homo sapien
```

```
        <400> 139
gaattcggca cgaggttccg tgcagctcac ggagaagcga atggacaaag tcggcaagta      60
ccccaaggag ctgcgcaagt gctgcgagga cggcatgcgg gagaacccca tgaggttctc     120
gtgccagcgc cggacccgtt tcatctccct ggcggaggcgt gcaagaaggt cttcctggac     180
tgctgcaact acatcacaga gctgcggcgg cagcacgcgc gggccagcca cctggcctgc     240
caggagtaac ctggatgagg acatcattgc agaagagaac atcgtttccc gaagtgagtt     300
cccagagagc tggctgtgga cgttgagga cttgaaagag ccaccgaaaa atggaatctc     360
tacgaagctc atgaatatat ttttgaaaga ctccatcacc acgtgggaga ttctggctgt     420
gagcatgtcg gacaagaaag ggatctgtgt ggcagacccc ttcgaggtca cagtaatgca     480
ggacttcttc atcgacctgc ggctaccctа                                     510
```

```
        <210> 140
        <211> 360
        <212> DNA
        <213> homo sapien
```

```
        <400> 140
gaattcggca cgagcggtaa ctaccccggc tgcgcacagc tcggcgctcc ttcccgctcc        60
ctcacacacc ggcctcagcc cgcaccggca gtagaagatg gtgaaagaaa caacttacta       120
cgatgttttg ggggtcaaac ccaatgctac tcaggaagaa ttgaaaaagg cttataggaa       180
actggctttg aagtaccatc ctgataagaa cccaaatgaa ggagagaagt ttaaacagat       240
ttctcaagct tacgaagttc tctctgatgc aaagaaaagg gaattatatg acaaaggagg       300
agaacaggca attaaagagg gtggagcagg tggcggtttt ggctccccca tggacatctt       360
```

```
        <210> 141
        <211> 483
        <212> DNA
        <213> homo sapien
```

```
        <400> 141
gaattcggca cgagagcaga ggctgatctt tgctggaaaa cagctggaag atgggctgca        60
ccctgtctga ctacaacatc cagaaagagt ccaccctgca cctggtgctc cgtctcagag       120
gtgggatgca aatcttcgtg aagacactca ctggcaagac catcaccctt gaggtggagc       180
ccagtgacac catcgagaac gtcaaagcaa agatccagga caaggaaggc attcctcctg       240
accagcagag gttgatcttt gccggaaagc agctggaaga tgggcgcacc ctgtctgact       300
acaacatcca gaaagagtct accctgcacc tggtgctccg tctcagaggt gggatgcaga       360
tcttcgtgaa gacccctgact ggtaagacca tcaccctcga ggtggagccc agtgacacca       420
tcgagaatgt caaggcaaag atccaagata ggaaggcat tcctcctgat cagcagaggt       480
tga                                                                     483
```

```
        <210> 142
        <211> 500
        <212> DNA
        <213> homo sapien
```

```
        <400> 142
gaattcggca cgaggcggcg acgaccgccg ggagcgtgtg cagcggcggc ggcggaagtg        60
gccggcgagc ccggtccccg ccggcaccat gcttcccttg tcactgctga agacggctca       120
gaatcacccc atgttggtgg agctgaaaaa tggggagacg tacaatggac acctggtgag       180
ctgcgacaac tggatgaaca ttaacctgcg agaagtcatc tgcacgtcca gggacgggga       240
caagttctgg cggatgcccg agtgctacat ccgcggcagc accatcaagt acctgcgcat       300
ccccgacgag atcatcgaca tggtcaagga ggaggtggtg gccaagggcc gcggccgcgg       360
aggcctgcag cagcagaagc agcagaaagg ccgcggcatg ggcggcgctg ccgaggtgt       420
gtttggtggc cggggccgag gtgggatccc gggcacaggc agaagccagc cagagaagaa       480
gcctggcaga caggcgggca                                                   500
```

```
        <210> 143
        <211> 400
        <212> DNA
        <213> homo sapien
```

```
        <400> 143
gaattcggca cgagctcgga tgtcagcagg cgtcccaacc cagcaggaac tggctcaatt        60
ctcagaagaa agcgatcggc cccgaggcag gaaggccggc tccggtgcag ggcgcgccgc       120
ctgcgggctg cttcgggcca gggtcgaccc gagggccagc gcaagcagcg gcaacaggag       180
cgccaggagg acatgaggct ctgcctgcag tcagcaactt ggaatattca gacttcagac       240
cagcatcaca gattataacc ctccgtaaat catctgcatc ccagctccca tcaaaagcca       300
gcctgaagga cccatggaca cgtgactcca gtgttctcaa caacatctta gatcaagttg       360
gtttgcacaa catttgcatc tacttgggac aaagcaagaa                             400
```

```
        <210> 144
```

```
        <211> 243
        <212> DNA
        <213> homo sapien

        <400> 144
gaattcggca cgagccagct cctaaccgcg agtgatccgc cagcctccgc ctcccgaggt      60
gcccggattg cagacggagt ctccttcact cagtgctcaa tggtgcccag gctggagtgc     120
agtggtgtga tctcggctcg ctacaacatc cacctcccag cagcctgcct tggcctccca     180
aagtgccgag attgcagcct ctgcccggcc gtcaccccgt ctgggaagtg aggagcgttt     240
ctg                                                                    243


        <210> 145
        <211> 450
        <212> DNA
        <213> homo sapien

        <400> 145
gaattcggca cgaggacagc aggaccgtgg aggccgcggc aggggtggca gtggtggcgg      60
cggcggcggc ggcggtggtg gttacaaccg cagcagtggt ggctatgaac ccagaggtcg     120
tggaggtggc cgtggaggca gaggtggcat gggcggaagt gaccgtggtg gcttcaataa     180
atttggtggc cctcgggacc aaggatcacg tcatgactcc gaacaggata attcagacaa     240
caacaccatc tttgtgcaag gcctgggtga gaatgttaca attgagtctg tggctgatta     300
cttcaagcag attggtatta ttaagacaaa caagaaaacg ggacagccca tgattaattt     360
gtacacagac agggaaactg gcaagctgaa gggagaggca acggtctctt ttgatgaccc     420
accttcagct aaagcagcct attgactggt                                      450


        <210> 146
        <211> 451
        <212> DNA
        <213> homo sapien

        <400> 146
gaattcggca cgagccatcg agtccctgcc tttcgacttg cagagaaatg tctcgctgat      60
gcgggagatc gacgcgaaat accaagagat cctgaaggag ctagacgagt gctacgagcg     120
cttcagtcgc gagacagacg gggcgcagaa gcggcggatg ctgcactgtg tgcagcgcgc     180
gctgatccgc accaggagct gggcgacgag aagatccaga tcgtgagcca gatggtggag     240
ctggtggaga accgcacgcg gcaggtggac agccacgtgg agctgttcga ggcgcagcag     300
gagctgggcg acacagcggg caacagcggc aaggctggcg cggacaggcc caaaggcgag     360
gcggcagcgc aggctgacaa gcccaacagc aagcgctcac ggcggcagcg caacaacgag     420
aaccgtgaga acgcgtccag caaccacgac c                                    451


        <210> 147
        <211> 400
        <212> DNA
        <213> homo sapien

        <400> 147
gaattcggca cgagctcgga tgtcagcagg cgtcccaacc cagcaggaac tggctcaatt      60
ctcagaagaa agcgatcggc cccgaggcag gaaggccggc tccggtgcag ggcgcgccgc     120
ctgcgggctg cttcgggcca gggtcgaccc gaggccagc gcaagcagcg gcaacaggag      180
cgccaggagg acatgaggct ctgcctgcag tcagcaactt ggaatattca gacttcagac     240
cagcatcaca gattataacc ctccgtaaat catctgcatc ccagctccca tcaaaagcca     300
gcctgaagga cccatggaca cgtgactcca gtgttctcaa caacatctta gatcaagttg     360
gtttgcacaa catttgcatc tacttgggac aaagcaagaa                           400
```

```
<210> 148
<211> 503
<212> DNA
<213> Homo sapien


<400> 148
aaaagaattc ggcacgagcg gcgccgctca tccccctctc ccagcagatt cccactggaa        60
attcgttgta tgaatcttat tacaagcagg tcgatccggc atacacaggg agggtggggg       120
cgagtgaagc tgcgcttttt ctaaagaagt ctggcctctc ggacattatc cttgggaaga       180
tatgggactt ggccgatcca gaaggtaaag ggttcttgga caaacagggt ttctatgttg       240
cactgagact ggtggcctgt gcacagagtg gccatgaagt taccttgagc aatctgaatt       300
tgagcatgcc accgcctaaa tttcacgaca ccagcagccc tctgatggtc acaccgccct       360
ctgcagaggc ccactgggct gtgagggtgg aagaaaaggc caaatttgat gggatttttg       420
aaagcctctt gcccatcaat ggtttgctct ctggagacaa agtcaagcca gtcctcatga       480
actcaaagct gcctcttgat gtc                                             503


<210> 149
<211> 1061
<212> DNA
<213> homo sapien


<400> 149
gaattcggca cgaggccttt tccagcaacc ccaaggtcca ggtggaggcc atcgaagggg        60
gagccctgca gaagctgctg gtcatcctgg ccacggagca gccgctcact gcaaagaaga       120
aggtcctgtt tgcactgtgc tccctgctgc gccacttccc ctatgcccag cggcagttcc       180
tgaagctcgg ggggctgcag gtcctgagga ccctggtgca ggagaagggc acggaggtgc       240
tcgccgtgcg cgtggtcaca ctgctctacg acctggtcac ggagaagatg ttcgccgagg       300
aggaggctga gctgacccag gagatgtccc cagagaagct gcagcagtat cgccaggtac       360
acctcctgcc aggcctgtgg gaacagggct ggtgcgagat cacgcccac ctcctggcgc       420
tgcccgagca tgatgcccgt gagaaggtgc tgcagacact gggcgtcctc ctgaccacct       480
gccgggaccg ctaccgtcag gaccccccagc tcggcaggac actggccagc ctgcaggctg       540
agtaccaggt gctggccagc ctggagctgc aggatggtga ggacgagggc tacttccagg       600
agctgctggg ctctgtcaac agcttgctga aggagctgag atgaggcccc acaccagtac       660
tggactggga tgccgctagt gaggctgagg ggtgccagcg tgggtgggct tctcaggcag       720
gaggacatct tggcagtgct ggcttggcca ttaaatggaa acctgaaggc catcctcttt       780
ctgctgtgtg tctgtgtaga ctgggcacag ccctgtggcc gggggggtcag gtgagtggtt       840
gggtgatggg ctctgctgac gtgcagggct cagcccaggg catccaggaa caggctccag       900
ggcaggaacc tgggcccagg agttgcaagt ctctgcttct taccaagcag cagctctgta       960
ccttgggaag tcgcttaatt gctctgagct tgtttcctca tctgtcagga gtgccattaa      1020
aggagaaaaa tcacgtaaaa aaaaaaaaaa aaaaactcga g                          1061


<210> 150
<211> 781
<212> DNA
<213> homo sapien


<400> 150
gaattcggca cgagaaatgg cggcaggggt cgaagcggca gccgaagtgg cggcgacaga        60
acccaaaatg gaggaagaga gcggcgcgcc ctgcgtgccg agcggcaacg gagctccggg       120
cccgaagggt gaagaacgac ctactcagaa tgagaagagg aaggagaaaa acataaaaag       180
aggaggcaat cgctttgagc catattccaa cccaactaaa agatacagag ccttcattac       240
aaatatacct tttgatgtga aatggcagtc acttaaagac ctggttaaag aaaaagttgg       300
tgaggtaaca tacgtggagc tcttaatgga cgctgaagga agtcaaggg gatgtgctgt       360
tgttgaattc aagatggagg agagcatgaa aaaagctgct gaagttctaa acaagcatag       420
tctgagtgga aggccactga aagtcaagga agatcctgat ggtgaacatg caaggagagc       480
```

```
aatgcaaaag gctggaagac ttggaagcac agtatttgta gcaaatctgg attataaagt    540
tggctggaag aaactgaagg aagtatttag tatggctggt gtggtggtcc gagcagacat    600
tctggaagat aaagatggga aaagtcgtgg aataggcatt gtgacttttg aacagtccat    660
tgaagctgtg caagcaatat ctatgtttaa tggccagttg ctgtttgata gaccgatgca    720
cgtcaagatg gatgagaggg cttttaccaaa gggagacttt tttcctcctg aacgccacag    780
c                                                                     781


        <210> 151
        <211> 3275
        <212> DNA
        <213> Homo sapien


        <400> 151
cttaagtgga tcctgcatca ggagggagca gacaccggag aaagaaaaac aagttgtgct     60
gtttgaggaa gcaagttgga cctgcactcc agcctgtgga gatgaaccta ggactgtgat    120
tctgctatcc agtatgttgg ctgaccacag gctcaaactg gaggattata aggatcgcct    180
gaaaagtgga gagcatctta atccagacca gttggaagct gtagagaaat atgaagaagt    240
gctacataat ttggaatttg ccaaggagct tcaaaaaacc ttttctgggt tgagcctaga    300
tctactaaaa gcgcaaaaga aggcccagag aagggagcac atgctaaaac ttgaggctga    360
gaagaaaaag cttcgaacta tacttcaagt tcagtatgta ttgcagaact tgacacagga    420
gcacgtacaa aaagacttca aaggggggttt gaatggtgca gtgtatttgc cttcaaaaga    480
acttgactac ctcattaagt tttcaaaact gacctgccct gaaagaaatg aaagtctgag    540
acaaacactt gaaggatcta ctgtctaaat tgctgaactc aggctatttt gaaagtatcc    600
cagttcccaa aaatgccaag gaaaaggaag taccactgga ggaagaaatg ctaatacaat    660
cagagaaaaa aacacaatta tcgaagactg aatctgtcaa agagtcagag tctctaatgg    720
aatttgccca gccagagata caaccacaag agtttcttaa cagacgctat atgacagaag    780
tagattattc aaacaaacaa ggcgaagagc aaccttggga agcagattat gctagaaaac    840
caaatctccc aaaacgttgg gatatgctta ctgaaccaga tggtcaagag aagaaacagg    900
agtcctttaa gtcctgggag gcttctggta agcaccagga ggtatccaag cctgcagttt    960
ccttagaaca gaggaaaaca gacacctcaa aactcaggtc tactctgccg gaagagcaga   1020
agaagcagga gatctccaaa tccaagccat ctcctagcca gtggaagcaa gatacaccta   1080
aatccaaagc agggtatgtt caagaggaac aaaagaaaca ggagacacca aagctgtggc   1140
cagttcagct gcagaaagaa caagatccaa agaagcaaac tccaaagtct tggacacctt   1200
ccatgcagag cgaacagaac accaccaagt catggaccac tcccatgtgt gaagaacagg   1260
attcaaaaca gccagagact ccaaaatcct gggaaaacaa tgttgagagt caaaaacact   1320
ctttaacatc acagtcacag atttctccaa agtcctgggg agtagctaca gcaagcctca   1380
taccaaatga ccagctgctg cccaggaagt tgaacacaga acccaaagat gtgcctaagc   1440
ctgtgcatca gcctgtaggt tcttcctcta cccttccgaa ggatccagta ttgaggaaag   1500
aaaaactgca ggatctgatg actcagattc aaggaacttg taactttatg caagagtctg   1560
ttcttgactt tgacaaacct tcaagtgcaa ttccaacgtc acaaccgcct tcagctactc   1620
caggtagccc cgtagcatct aaagaacaaa atctgtccag tcaaagtgat tttcttcaag   1680
agccgttaca ggtatttaac gttaatgcac ctctgcctcc acgaaaagaa caagaaataa   1740
aagaatcccc ttattcacct ggctacaatc aaagtttttac cacagcaagt acacaaacac   1800
cacccagtg ccaactgcca tctatacatg tagaacaaac tgtccattct caagagactg    1860
cagcaaatta tcatcctgat ggaactattc aagtaagcaa tggtagcctt gcctttttacc   1920
cagcacagac gaatgtgttt cccagaccta ctcagccatt tgtcaatagc cggggatctg   1980
ttagaggatg tactcgtggt gggagattaa taaccaattc ctatcggtcc cctggtggtt   2040
ataaaggttt tgatacttat agaggactcc cttcaatttc caatggaaat tatagccagc   2100
tgcagttcca agctagagag tattctggag caccttattc ccaaagggat aatttccagc   2160
agtgttataa gcgaggaggg acatctggtg gtccacgagc aaattcgaga gcagggtgga   2220
gtgattcttc tcaggtgagc agcccagaaa gagacaacga aacctttaac agtggtgact   2280
ctggacaagg agactcccgt agcatgaccc ctgtggatgt gccagtgaca aatccagcag   2340
ccaccatact gccagtacac gtctaccctc tgcctcagca gatgcgagtt gccttctcag   2400
cagccagaac ctctaatctg gcccctggaa cttttagacca acctattgtg tttgatcttc   2460
ttctgaacaa cttaggagaa acttttgatc ttcagcttgg tagatttaat tgcccagtga   2520
```

```
atggcactta cgttttcatt tttcacatgc taaagctggc agtgaatgtg ccactgtatg    2580
tcaacctcat gaagaatgaa gaggtcttgg tatcagccta tgccaatgat ggtgctccag    2640
accatgaaac tgctagcaat catgcaattc ttcagctctt ccagggagac cagatatggt    2700
tacgtctgca caggggagca atttatggaa gtagctggaa atattctacg ttttcaggct    2760
atcttcttta tcaagattga aagtcagtac agtattgaca ataaaaggat ggtgttctaa    2820
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ttttgttcct agagggagga ggtccttact tttttgtttt ccttcctgag gtgaaaaatc    2940
aagctgaatg acaattagca ctaatctggc actttataaa ttgtgatgta gcctcgctag    3000
tcaagctgtg aatgtatatt gtttgcactt aatccttaac tgtattaacg ttcagcttac    3060
taaactgact gcctcaagtc caggcaagtt acaatgcctt gttgtgcctc aataaaaaag    3120
ttacatgcaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa    3180
aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa    3240
aaaaaaaaaa aaaaaaaaaa aaaaaaaaac tcgag                               3275
```

```
<210> 152
<211> 2179
<212> DNA
<213> homo sapien

<400> 152
gaattcggca ccaggcacta ttaaatgtga ggcagcctcc atctactaca acatttgtgc      60
tgaatcaaat aaatcatctt ccacccttgg gatctacaat tgtaatgact aaaacaccac     120
ctgtaacaac caacaggcaa accatcactt taactaagtt tatccagact actgcaagca     180
cacgcccgtc agtctcagca ccaacagtac gaaatgccat gacctctgca ccttcaaaag     240
accaagttca gcttaaagat ctactgaaaa ataatagtct taatgaactg atgaaactaa     300
agccacctgc taatattgct cagccagtag caacagcagc tactgatgta agcaatggta     360
cagtaaagaa agagtcttct aataaagaag gagctagaat gtggataaac gacatgaaga     420
tgaggagttt ttccccaacc atgaaggttc ctgttgtaaa agaagatgat gaaccagagg     480
aagaagatga agaagaaatg ggtcatgcag aaacctatgc agaatacatg ccaataaaat     540
taaaaattgg cctacgtcat ccagatgctg tagtggaaac cagctctta tccagtgtta     600
ctcctcctga tgtttggtac aaaacatcca tttctgagga aaccattgat aatggctggt     660
tatcagcatt gcagcttgag gcaattacat atgcagccca gcaacatgaa actttcctac     720
ctaatggaga tcgtgctggc ttcttaatag gtgatggtgc cggtgtagga aaaggaagga     780
cgatagcagg aatcatctat gaaaattatt tgttgagtag aaaacgagca ttgtggttta     840
gtgtttcaaa tgacttaaag tatgatgctg aaagagattc aagggatatt ggagcaaaaa     900
acattttggt tcattcgtta aataagttta aatacggaaa aatttcttcc aaacataatg     960
ggagtgtgaa aaagggtgtt atttttgcta cttactcttc acttattggt gaaagccagt    1020
ctggcggcaa gtataaaact aggttaaaac aacttctgca ttggtgcggt gatgacttcg    1080
atggagtgat agtgtttgat gagtgtcata agccaaaaa cttatgtcct gttggttctt    1140
caaagccaac caagacaggc ttagcagttt tagagcttca gaacaaattg ccaaaagcca    1200
gagttgttta tgctagtgca actggtgctt ctgaaccacg caacatggcc tatatgaacc    1260
gtcttggcat atggggtgag ggtactccat ttagagaatt cagtgatttt attcaagcag    1320
tagaacggag aggagttggt gccatggaaa tagttgctat ggatatgaag cttagaggaa    1380
tgtacattgc tcgacaactg agctttactg gagtgacctt caaaattgag gaagttcttc    1440
tttctcagag ctacgttaaa atgtataaca aagctgtcaa gctgtgggtc attgccagag    1500
agcggtttca gcaagctgca gatctgattg atgctgagca acgaatgaag aagtccatgt    1560
ggggtcagtt ctggtctgct caccagaggt cttcaaaata cttatgcata gcatccaaag    1620
ttaaaagggt tgtgcaacta gctcgagagg aaatcaagaa tggaaaatgt gttgtaattg    1680
gtctgcagtc tacaggagaa gctagaacat tagaagcttt ggaagagggc ggggggagaat    1740
tgaatgattt tgtttcaact gccaaaggtg tgttgcagtc actcattgaa aaacattttc    1800
ctgctccaga caggaaaaaa cttatagtt tactaggaat cgatttgaca gctccaagta    1860
acaacagttc gccaagagat agtccttgta aagaaataa aataaagaag cggaaaggtg    1920
aagaaataac tcgagaagcc aaaaaagcac gaaaagtagg tggccttact ggtagcagtt    1980
ctgacgacag tggaagtgaa tctgatgcct ctgataatga agaaagtgac tatgagagct    2040
ctaaaaacat gagttctgga gatgatgacg atttcaaccc attttttagat gagtctaatg    2100
```

aggatgatga aaatgatccc tggttaatta aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa   2160
aaaaaaaaaa aaactcgag   2179

        <210> 153
        <211> 2109
        <212> DNA
        <213> Homo sapien

        <400> 153
cagagagccc caggcatcga ggagaaggcg gcggagaatg gggccctggg gtcccccgag   60
agagaagaga aagtgctgga gaatggggag ctgacacccc caaggaggga ggagaaagcg   120
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cccgagagtt ggggtccagc ccccacgatc ggggagccag ccccagagac ctctctggag   360
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cccgagggag agcccggggc cccagacagc agggccggcg gagacacggc actcagcgga   780
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tctccccttt ccctccccag cccggccct ggaggggcct ctggttcaaa ccttcgcgtg   2040
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aaactcgag   2109

        <210> 154
        <211> 1411
        <212> DNA
        <213> homo sapien

        <400> 154
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cccaagacct gatacaaata aggatgtata caaagtattg ccagaatcca agaaggcacc   180

```
gggcagtggt gcagtatttg agaggaacgg accacatgct agcagtagtg gggtgctccc    240
tttgggactc cagcctgcgc ctggactttc caagtcacta tcctctcagg tgtggcaacc    300
aagtcctgac ccttggcatc ctggagaaca atcctgtgaa ctcagtactt gtcgacagca    360
gttggaattg atccgtttac agatggagca aatgcagctt cagaacggag ccatgtgtca    420
ccatcctgct gctttcgctc cattactgcc cacectagag ccagcacagt ggctcagcat    480
cctgaacagt aacgagcatc tcctgaagga gaaggagctc ctcattgaca agcaaaggaa    540
gcatatctct cagctggagc agaaagtgcg agagagtgaa ctgcaagtcc acagtgccct    600
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agagaacact ttcttacggg cacagtttgc acagaagaca gaagccctga gcaaggagaa    720
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aggtcgtgat aaacatatca ataatttgaa aaagaaatgt cagaaggaat cagagcagaa    900
ccggagaag cagcagcgta ttgaaacctt ggagcgctat ctagctgacc tgcccaccct    960
agaagaccat cagaaacaga cggagcagct taaggacgct gaattaaaga acacagaact   1020
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aattgaaagt cacttatggc cgggtgtggt ggcgcacacc tttaatccca gcacttggga   1320
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```

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        <210> 155
        <211> 678
        <212> DNA
        <213> homo sapien

        <400> 155
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cccgcccctg tgttggtccc acaatccctg gcaatgagag gccagggttt attggacaga    180
gtcagttgtg gggttcagag ggtcagcaat caatcaatcc tccgaatcca gagatttaga    240
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ggaacctgtc ttttgatttg gggttcaaca tacagagttc aggtacctgc aggaatttgc    360
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ggaagcccac cgacgccagc gtcacccttt cttactcctt ggcactacgg ctaacaggac    540
gcaatccctc aactacggct gcatcgttga gaatccacag acacacgagg tattgcacta    600
tgtggagaaa cccagcacat ttatcagtga catcatcaac tgcggcacct acctcttttc    660
tcctgaagcc ttgaagcc                                                  678
```

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        <210> 156
        <211> 2668
        <212> DNA
        <213> Homo sapien

        <400> 156
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tgccccggccc aggcgacctg cggagcgcaa cgaggccgcg ggcggccgag ggctggtcgg    180
cgcccatcct gacccctggca cgcagggcca ccgggaacct gtcggcgagc tgcgggagcg    240
cgctgcgcgc ggccgcgggg ctgggcggcg gggacagcgg ggacggcacg gcgcgcgcag    300
cttctaagtg ccagatgatg gaggagcgtg ccaacctgat gcacatgatg aaactcagca    360
tcaaggtgtt gctccagtcg gctctgagcc tgggccgcag cctggatgcg gaccatgccc    420
ccttgcagca gttctttgta gtgatggagc actgcctcaa acatgggctg aaagttaaga    480
```

```
agagttttat tggccaaaat aaatcattct ttggtccttt ggagctggtg gagaaacttt        540
gtccagaagc atcagatata gcgactagtg tcagaaatct tccagaatta aagacagctg        600
tgggaagagg ccgagcgtgg ctttatcttg cactcatgca aaagaaactg gcagattatc        660
tgaaagtgct tatagacaat aaacatctct taagcgagtt ctatgagcct gaggctttaa        720
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cagctgcaac agaccgaatt tgctcacttc aagaagaaca gcagcagtta agagaacaaa       1080
atgaattaat tcgagaaaga agtgaaaaga gtgtagagat aacaaaacag gataccaaag       1140
ttgagctgga gacttacaag caaactcggc aaggtctgga tgaaatgtac agtgatgtgt       1200
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aagcgacaca ctgtaggcag tgtgagaagg agttctccat ttcccggaga aagcaccact       1980
gccggaactg tggccacatc ttctgcaaca cctgctccag caacgagctg gccctgccct       2040
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tcaaagaatt taacctattt tacagcagtt cagttctgct agtgagtagt tttcctctcc       2460
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gtgccttctc tgcttcagac aagagatctg ccatttcatg cccttgtgac tacctatcat       2580
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aaaaaaaaaa aaaaaaaaaa aactcgag                                         2668
```

<210> 157
<211> 2313
<212> DNA
<213> homo sapien

<400> 157

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ccagctcaag gtgctgtccc acaacctgta cacggtcctg cacatccccc atgaccccgt        180
ggccctggag gaacacttcc gagatgatga tgacggccct gtgtccagcc agggatacat        240
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ggcccaggtg gcccagacca ccgggggggct cagcgtctgg cagttcctgg agctcttcaa        600
ttcgggccgc tgcctgcggg gcgtggggccg ggacaccctc agcatggcca tccacgaggt        660
```

```
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gtatgagatg agcgcctcag acacgcgcca gcgccaggag tggacagctg ccatccagat    960
ggcgatccgg ctgcaggccg aggggaagac gtccctacac aaggacctga agcagaaacg   1020
gcgcgagcag cgggagcagc gggagcggcg ccgggcggcc aaggaagagg agctgctgcg   1080
gctgcagcag ctgcaggagg agaaggagcg gaagctgcag gagctggagc tgctgcagga   1140
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ccagctccaa gccccagacc atgggagctg tctgggatgt tgatccttga gaacttggcc   2160
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ggctacctgc ccccaggtgg ccaccaagtt gtggaagcac atttctaaat aaaaactgct   2280
cttagaatga aaaaaaaaaa aaaaaactc gag                                 2313
```

<210> 158
<211> 2114
<212> DNA
<213> homo sapien

<400> 158

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gacacctcag cagcgcacca ggtggtttta ggagaaaact tgatagccac agccctttgt   180
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caggatgtga caataaagcg ggaagaaaca aagaagaaga tagagaaaga gaagaaggag   780
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cagttagtga ggaacggagc caagctgagc agccttcctc aaatccctac tcccacttta  1200
```

```
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tggttgtggt agtcggcact ttggtagtgt gagctgttcc tggggtggaa gaggggtgg    1740
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agaagggagt ggctttggag ctggtgactg tgggtgtcgt ggccgtggtg ctcacatgtg   2040
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```

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    <210> 159
    <211> 278
    <212> DNA
    <213> homo sapien


    <400> 159
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tgcagaatga gaatcactcc taaaataggt aatggtaaaa attaaattga caattacctc   180
tctctatgca gaaggaaata tcacctatat gacatcatca tcatctattg atacttgctg   240
gcagtgctaa taatggtttt aatgccaatt tgtaagaa                           278
```

```
    <210> 160
    <211> 848
    <212> DNA
    <213> homo sapien


    <400> 160
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agccaggagc ttggggaagg gaagcgcgcc cccggggccg gtcccggagg gctcgatccg   120
catctacagc atgaggttct gcccgtttgc tgagaggacg cgtctagtcc tgaaggccaa   180
gggaatcagg catgaagtca tcaatatcaa cctgaaaaat aagcctgagt ggttctttaa   240
gaaaaatccc tttggtctgg tgccagttct ggaaaacagt cagggtcagc tgatctacga   300
gtctgccatc acctgtgagt acctggatga agcatacc ca gggaagaagc tgttgccgga   360
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cttggtagga agctttatta gaagccaaaa taaagaagac tatgctggcc taaaagaaga   480
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cctagagctc tacttacaga acagccctga ggcctgtgac tatgggctct gaaggggggca   780
ggagtcagca ataaagctat gtctgatatt ttccttcact aaaaaaaaaa aaaaaaaaaa   840
aactcgag                                                             848
```

```
    <210> 161
    <211> 432
    <212> DNA
    <213> homo sapien
```

<400> 161
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ggaagggaag ggcccaagta aagcacagcg cgggagccta gagcacatga agctgatcct   180
gcgtgataag gagaaggagg tggaatgtca gcaggagcat atccatgaac tccaggagct   240
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cctcctgtcc cagcgagagc aggaaatagt ggtcctgcag cagcaactgc aggaagccag   360
ggaacaaggg gagctgaagg agcagtcact tcagagtcaa ctggatgagg cccagagagc   420
cctagcccag ag   432

    <210> 162
    <211> 433
    <212> DNA
    <213> homo sapien

    <400> 162
gattcggcac gagccggagc tgggttgctc ctgctcccgt ctccaagtcc tggtacctcc   60
ttcaagctgg gagagggctc tagtccctgg ttctgaacac tctggggttc tcgggtgcag   120
gccgccatga gcaaacggaa ggcgccgcag gagactctca acgggggaat caccgacatg   180
ctcacagaac tcgcaaactt tgagaagaac gtgagccaag ctatccacaa gtacaatgct   240
tacagaaaag cagcatctgt tatagcaaaa tacccacaca aaataaagag tggagctgaa   300
gctaagaaat tgcctggagt aggaacaaaa attgctgaaa agattgatga gtttttagca   360
actggaaaat tacgtaaact ggaaaagatt cggcaggatg atacgagttc atccatcaat   420
ttcctgactc gag   433

    <210> 163
    <211> 432
    <212> DNA
    <213> homo sapien

    <400> 163
gaattcggca ccagatgagg ccaacgaggt gacggacagc gcgtacatgg gctccgagag   60
cacctacagt gagtgtgaga ccttcacgga cgaggacacc agcaccctgg tgcaccctga   120
gctgcaacct gaaggggacg cagacagtgc cggcggctcg gccgtgccct ctgagtgcct   180
ggacgccatg gaggagcccg accatggtgc cctgctgctg ctcccaggca ggcctcaccc   240
ccatggccag tctgtcatca cggtgatcgg gggcgaggag cactttgagg actacggtga   300
aggcagtgag gcggagctgt ccccagagac cctatgcaac gggcagctgg gctgcagtga   360
ccccgctttc ctcacgccca gtccgacaaa gcggctctcc agcaagaagg tggcaaggta   420
cctgcaccag tc   432

    <210> 164
    <211> 395
    <212> DNA
    <213> homo sapien

    <400> 164
gacacttgaa tcatgggtga cgttaaaaat tttctgtatg cctggtgtgg caaaaggaag   60
atgaccccat cctatgaaat tagagcagtg gggaacaaaa acaggcagaa attcatgtgt   120
gaggttcagg tggaaggtta taattacact ggcatgggaa attccaccaa taaaaaagat   180
gcacaaagca atgctgccag agactttgtt aactatttgg ttcgaataaa tgaaataaag   240
agtgaagaag ttccagcttt tggggtagca tctccgcccc cacttactga tactcctgac   300
actacagcaa atgctgaagg catcttgttg acatcgaata tgactttgat aataaatacc   360
ggttcctgaa aaaaaaaaaa aaaaaaaaac tcgag   395

```
        <210> 165
        <211> 503
        <212> DNA
        <213> homo sapien

        <400> 165
gaattcggca ccaggaacgc tcggtgagag gcggaggagc ggtaactacc ccggttgcgc     60
acagctcggc gctccttccc gctccctcac acaccggcct cagcccgcac cggcagtaga    120
agatggtgaa agaaacaact tactacgatg ttttgggggt caaacccaat gctactcagg    180
aagaattgaa aaaggcttat aggaaactgg ccttgaagta ccatcctgat aagaacccaa    240
atgaaggaga gaagtttaaa cagatttctc aagcttacga agttctctct gatgcaaaga    300
aaagggaatt atatgacaaa ggaggagaac aggcaattaa agagggtgga gcaggtggcg    360
gttttggctc ccccatggac atctttgata tgttttttgg aggaggagga aggatgcaga    420
gagaaaggag aggtaaaaat gttgtacatc agctctcagt aaccctagaa gacttatata    480
atggtgcaac aagaaaactg gct                                           503


        <210> 166
        <211> 893
        <212> DNA
        <213> homo sapien

        <400> 166
gaattcggca cgagaggaac ttctcttgac gagaagagag accaaggagg ccaagcaggg     60
gctgggccag aggtgccaac atggggaaac tgaggctcgg ctcggaaggg tgagagtgag    120
actacatctc aaaaaaaaaa aaaaaaaaaa aaagaaaga aaagaaaaga aaaagaaag    180
aacggaagta gttgtaggta gtggtatggt ggtatgagtc tgttttctgt tacttataac    240
aacaacaaca acaaaaaacg ctgaaactgg gtaatttata aagaaaagga aaaaaagcag    300
aaaaaaatca ggaagaagag aaaggaaaag aagacaaata aatgaaattt atgtattaca    360
gttctgaagg ctgagacatc ccaggtcaag ggtccacact tggcgagggc tttcttgctg    420
gtggagactc tttgtggagt cctgggacag tgcagaagga tcacgcctcc ctaccgctcc    480
aagcccagcc ctcagccatg gcatgccccc tggatcaggc cattggcctc ctcgtggcca    540
tcttccacaa gtactccggc agggagggtg acaagcacac cctgagcaag aaggagctga    600
aggagctgat ccagaaggag ctcaccattg gctcgaagct gcaggatgct gaaattgcaa    660
ggctgatgga agacttggac cggaacaagg accaggaggt gaacttccag gagtatgtca    720
ccttcctggg ggccttggct ttgatctaca atgaagccct caagggctga aaataaatag    780
ggaagatgga gacaccctct gggggtcctc tctgagtcaa atccagtggt gggtaattgt    840
acaataaatt tttttggtc aaatttaaaa aaaaaaaaa aaaaaactc gag            893


        <210> 167
        <211> 549
        <212> DNA
        <213> homo sapien

        <400> 167
gaattcggca cgagcccaga tcccgaggtc cgacagcgcc cggcccagat ccccacgcct     60
gccaggagca agccgagagc cagccggccg gcgcactccg actccgagca gtctctgtcc    120
ttcgacccga gccccgcgcc ctttccggga cccctgcccc gcgggcagcg ctgccaacct    180
gccggccatg gagacccgt cccagcggcg cgccacccgc agcggggcgc aggccagctc    240
cactccgctg tcgcccaccc gcatcacccg gctgcaggag aaggaggacc tgcaggagct    300
caatgatcgc ttggcggtct acatcgaccg tgtgcgctcg ctggaaacgg agaacgcagg    360
gctgcgcctt cgcatcaccg agtctgaaga ggtggtcagc cgcgaggtgt ccggcatcaa    420
ggccgcctac gaggccgagc tcgggggatgc ccgcaagacc cttgactcag tagccaagga    480
gcgcgcccgc ctgcagctgg agctgagcaa agtgcgtgaa gagtttaagg agctgaaagc    540
gcgcaatac                                                           549
```

<210> 168
<211> 547
<212> DNA
<213> homo sapien

<400> 168

```
gaattcggca cgagatggcg gcaggggtcg aagcggcggc ggaggtggcg gcgacggaga     60
tcaaaatgga ggaagagagc ggcgcgcccg gcgtgccgag cggcaacggg gctccgggcc    120
ctaagggtga aggagaacga cctgctcaga atgagaagag gaaggagaaa aacataaaaa    180
gaggaggcaa tcgctttgag ccatatgcca atccaactaa aagatacaga gccttcatta    240
caaacatacc ttttgatgtg aaatggcagt cacttaaaga cctggttaaa gaaaaagttg    300
gtgaggtaac atacgtggag ctcttaatgg acgctgaagg aaagtcaagg ggatgtgctg    360
ttgttgaatt caagatggaa gagagcatga aaaaagctgc ggaagtccta aacaagcata    420
gtctgagcgg aagaccactg aaagtcaaag aagatcctga tggtgaacat gccaggagag    480
caatgcaaaa ggctggaaga cttggaagca cagtatttgt agcaaatctg gattataaag    540
ttggctg                                                              547
```

<210> 169
<211> 547
<212> DNA
<213> homo sapien

<400> 169

```
gaattcggca ccaggagtcc gactgtgctc gctgctcagc gccgcacccg gaagatgagg     60
ctcgccgtgg gagccctgct ggtctgcgcc gtcctggggc tgtgtctggc tgtccctgat    120
aaaactgtga gatggtgtgc agtgtcggag catgaggcca ctaagtgcca gagtttccgc    180
gaccatatga aaagcgtcat tccatccgat ggtcccagtg ttgcttgtgt gaagaaagcc    240
tcctaccttg attgcatcag ggccattgcg gcaaacgaag cggatgctgt gacactggat    300
gcaggtttgg tgtatgatgc ttacctggct cccaataacc tgaagcctgt ggtggcagag    360
ttctatgggt caaaagagga tccacagact ttctattatg ctgttgctgt ggtgaagaag    420
gatagtggct tccagatgaa ccagcttcga ggcaagaagt cctgccacac gggtctaggc    480
aggtccgctg ggtggaacat ccccataggc ttacttact gtgacttacc tgagccacgt    540
aaacctc                                                              547
```

<210> 170
<211> 838
<212> DNA
<213> homo sapien

<400> 170

```
gaattcggca ccagaggagc tcggcctgcg ctgcgccacg atgtccgggg agtcagccag     60
gagcttgggg aaggaagcg cgcccccggg gccggtcccg gagggctcga tccgcatcta    120
cagcatgagg ttctgcccgt ttgctgagag gacgcgtcta gtcctgaagg ccaagggaat    180
caggcatgaa gtcatcaata tcaacctgaa aaataagcct gagtggttct ttaagaaaaa    240
tcccttlggt ctggtgccag ttctggaaaa cagtcagggt cagctgatct acgagtctgc    300
catcacctgt gagtacctgg atgaagcata cccaggaaag aagctgttgc cggatgaccc    360
ctatgagaaa gcttgccaga agatgatctt agagttgttt tctaaggtgc catccttggt    420
aggaagcttt attagaagcc aaaataaaga agactatgat ggcctaaaag aagaatttcg    480
taaagaattt accaagctag aggaggttct gactaataag aagacgacct tctttggtgg    540
caattctatc tctatgattg attacctcat ctggccctgg tttgaacggc tggaagcaat    600
gaagttaaat gagtgtgtag accacactcc aaaactgaaa ctgtggatgg cagccatgaa    660
ggaagatccc acagtctcag ccctgcttac tagtgagaaa gactggcaag gtttcctaga    720
gctctactta cagaacagcc ctgaggcctg tgactatggg ctctgaaggg ggcaggagtc    780
agcaataaag ctatgtctga tattttcctt cactaaaaaa aaaaaaaaaa aactcgag     838
```

86-

```
<210> 171
<211> 547
<212> DNA
<213> homo sapien

<400> 171
gaattcggca ccagcgggat ttgggtcgca gttcttgttt gtggattgct gtgatcgtca      60
cctgacaatg cagatcttcg tgaagactct gactggtaag accatcaccc tcgaggttga     120
gcccagtgac accatcgaga atgtcaaggc aaagatccaa gataaggaag gcatccctcc     180
tgaccagcag aggctgatct ttgctggaaa acagctggaa gatgggcgca ccctgtctga     240
ctacaacatc cagaaagagt ccaccctgca cctggtgctc cgtctcagag gtgggatgca     300
aatcttcgtg aagacactca ctggcaagac catcaccctt gaggtcgagc ccagtgacac     360
catcgagaac gtcaaagcaa agatccagga caaggaaggc attcctcctg accagcagag     420
gttgatcttt gccggaaagc agctggaaga tgggcgcacc ctgtctgact acaacatcca     480
gaaagagtct accctgcacc tggtgctccg tctcagaggt gggatgcaga tcttcgtgaa     540
gaccctg                                                                547
```

```
<210> 172
<211> 608
<212> DNA
<213> homo sapien

<400> 172
gaattcggca ccagagactt ctccctctga ggcctgcgca cccctcctca tcagcctgtc      60
caccctcatc tacaatggtg ccctgccatg tcagtgcaac cctcaaggtt cactgagttc     120
tgagtgcaac cctcatggtg gtcagtgcct gtgcaagcct ggagtggttg ggcgccgctg     180
tgacctctgt gcccctggct actatggctt tggccccaca ggctgtcaag gcgcttgcct     240
gggctgccgt gatcacacag ggggtgagca ctgtgaaagg tgcattgctg gtttccacgg     300
ggacccacgg ctgccatatg ggggccagtg ccggccctgt ccctgtcctg aaggccctgg     360
gagccaacgg cactttgcta cttcttgcca ccaggatgaa tattcccagc agattgtgtg     420
ccactgccgg gcaggctata cggggctgcg atgtgaagct tgtgcccctg ggcactttgg     480
ggacccatca aggccaggtg gccggtgcca actgtgtgag tgcagtggga acattgatcc     540
aatggatcct gatgcctgtg accccacac ggggcaatgc ctgcgctgtt tacaccacac     600
agagggtc                                                               608
```

```
<210> 173
<211> 543
<212> DNA
<213> homo sapien

<400> 173
gaattcggca ccagagatca tccgccagca gggtctggcc tcctacgact acgtgcgccg      60
ccgcctcacg gctgaggacc tgttcgaggc tcggatcatc tctctcgaga cctacaacct     120
gctccgggag ggcaccagga gcctccgtga ggctctcgag gcggagtccg cctggtgcta     180
cctctatggc acgggctccg tggctggtgt ctacctgccc ggttccaggc agacactgag     240
catctaccag gctctcaaga aagggctgct gagtgccgag gtggcccgcc tgctgctgga     300
ggcacaggca gccacaggct tcctgctgga cccggtgaag ggggaacggc tgactgtgga     360
tgaagctgtg cggaagggcc tcgtggggcc cgaactgcac gaccgcctgc tctcggctga     420
gcgggcggtc accggctacc gtgaccccta caccgagcag accatctcgc tcttccaggc     480
catgaagaag gaactgatcc ctactgagga ggccctgcgg ctgtggatgc ccagctggcc     540
acc                                                                    543
```

```
<210> 174
<211> 548
<212> DNA
```

<213> homo sapien

<400> 174
```
gaattcggca cgagaaatgg cggcaggggt cgaagcggcg gcggaggtgg cggcgacgga       60
gatcaaaatg gaggaagaga gcggcgcgcc cggcgtgccg agcggcaacg gggctccggg      120
ccctaagggt gaaggagaac gacctgctca gaatgagaag aggaaggaga aaaacataaa      180
aagaggaggc aatcgctttg agccatatgc caatccaact aaaagataca gagccttcat      240
tacaaacata ccttttgatg tgaaatggca gtcacttaaa gacctggtta aagaaaaagt      300
tggtgaggta acatacgtgg agctcttaat ggacgctgaa ggaaagtcaa ggggatgtgc      360
tgttgttgaa ttcaagatgg aagagagcat gaaaaaagct gcggaagtcc taaacaagca      420
tagtctgagc ggaagaccac tgaaagtcaa agaagatcct gatggtgaac atgccaggag      480
agcaatgcaa aaggtgatgg ctacgactgg tgggatgggt atgggaccag gtggcccagg      540
aatgatta                                                                548
```

<210> 175
<211> 604
<212> DNA
<213> homo sapien

<400> 175
```
gaattcggca ccagaggacc tccaggacat gttcatcgtc cataccatcg aggagattga       60
gggcctgatc tcagcccatg accagttcaa gtccaccctg ccggacgccg atagggagcg      120
cgaggccatc ctggccatcc acaaggaggc ccagaggatc gctgagagca accacatcaa      180
gctgtcgggc agcaacccct acaccaccgt caccccgcaa atcatcaact ccaagtggga      240
gaaggtgcag cagctggtgc caaaacggga ccatgccctc ctggaggagc agagcaagca      300
gcagtccaac gagcacctgc gccgccagtt cgccagccag gccaatgttg tggggccctg      360
gatccagacc aagatggagg agatcgggcg catctccatt gagatgaacg ggaccctgga      420
ggaccagctg agccacctga agcagtatga acgcagcatc gtggactaca agcccaacct      480
ggacctgctg gagcagcagc accagcttat ccaggaggcc ctcatcttcg acaacaagca      540
caccaactat accatggagc acatccgcgt gggctgggag cagctgctca ccaccattgc      600
ccgg                                                                    604
```

<210> 176
<211> 486
<212> DNA
<213> homo sapien

<400> 176
```
gaattcggca ccagccaagc tcactattga atccacgccg ttcaatgtcg cagaggggaa       60
ggaggttctt ctactcgccc acaacctgcc ccagaatcgt attggttaca gctggtacaa      120
aggcgaaaga gtggatggca acagtctaat tgtaggatat gtaataggaa ctcaacaagc      180
tacccccaggg cccgcataca gtggtcgaga gacaatatac cccaatgcat ccctgctgat      240
ccagaacgtc acccagaatg acacaggatt ctataccta caagtcataa agtcagatct      300
tgtgaatgaa gaagcaaccg gacagttcca tgtataccccg gagctgccca gccctccat      360
ctccagcaac aactccaacc ccgtggagga caaggatgct gtggccttca cctgtgaacc      420
tgaggttcag aacacaacct acctgtggtg ggtaaatggt cagagcctcc cggtcagtcc      480
caaggc                                                                  486
```

<210> 177
<211> 387
<212> DNA
<213> homo sapien

<400> 177
```
gaattcggca ccagggacag cagaccagac agtcacagca gccttgacaa aacgttcctg       60
```

```
gaactcaagc tcttctccac agaggaggac agagcagaca gcagagacca tggagtctcc    120
ctcggcccct ccccacagat ggtgcatccc ctggcagagg ctcctgctca cagcctcact    180
tctaaccttc tggaacccgc ccaccactgc caagctcact attgaatcca cgccgttcaa    240
tgtcgcagag gggaaggagg tgcttctact tgtccacaat ctgccccagc atcttttgg     300
ctacagctgg tacaaaggtg aaagagtgga tggcaaccgt caaattatag gatatgtaat    360
aggaactcaa caagctaccc cagggcc                                        387
```

    <210> 178
    <211> 440
    <212> DNA
    <213> homo sapien

    <400> 178
```
gaattcggca cgaggagaag cagaaaaaca aggaatttag ccagacttta gaaaatgaga     60
aaaatacctt actgagtcag atatcaacaa aggatggtga actaaaaatg cttcaggagg    120
aagtaaccaa aatgaacctg ttaaatcagc aaatccaaga agaactctct agagttacca    180
aactaaagga gacagcagaa gaagagaaag atgatttgga agagaggctt atgaatcaat    240
tagcagaact taatggaagc attgggaatt actgtcagga tgttacagat gcccaaataa    300
aaaatgagct attggaatct gaaatgaaga accttaaaaa gtgtgtgagt gaattggaag    360
aagaaaagca gcagttagtc aaggaaaaaa ctaaggtgga atcagaaata cgaaaggaat    420
atttggagaa aatacaaggt                                                440
```

    <210> 179
    <211> 443
    <212> DNA
    <213> homo sapien

    <400> 179
```
gaattcggca ccagcggggg gctacggcgg cggctacggc ggcgtcctga ccgcgtccga     60
cgggctgctg gcgggcaacg agaagctaac catgcagaac ctcaacgacc gcctggcctc    120
ctacctggac aaggtgcgcg ccctggaggc ggccaacggc gagctagagg tgaagatccg    180
cgactggtac cagaagcagg ggcctgggcc ctcccgcgac tacagccact actacacgac    240
catccaggac ctgcgggaca agattcttgg tgccaccatt gagaactcca ggattgtcct    300
gcagatcgac aacgcccgtc tggctgcaga tgacttccga accaagtttg agacggaaca    360
ggctctgcgc atgagcgtgg aggccgacat caacggcctg cgcagggtgc tggatgagct    420
gaccctggcc aggaccgacc tgg                                            443
```

    <210> 180
    <211> 403
    <212> DNA
    <213> homo sapien

    <400> 180
```
gaattcggca cgaggttatg agagtcgact tcaatgttcc tatgaagaac aaccagataa     60
caaacaacca gaggattaag gctgctgtcc caagcatcaa attctgcttg gacaatggag    120
ccaagtcggt agtccttatg agccacctag gccggcctga tggtgtgccc atgcctgaca    180
agtactcctt agagccagtt gctgtagaac tcagatctct gctgggcaag gatgttctgt    240
tcttgaagga ctgtgtaggc ccagaagtgg agaaagcctg tgccaaccca gctgctgggt    300
ctgtcatcct gctggagaac ctccgctttc atgtggagga agaagggaag ggaaaagatg    360
cttctgggaa caaggttaaa gccgagccag ccaaaataga agc                      403
```

    <210> 181
    <211> 493
    <212> DNA
    <213> homo sapien

```
    <400> 181
gaattcggca ccagcagagg tctccagagc cttctctctc ctgtgcaaaa tggcaactct      60
taaggaaaaa ctcattgcac cagttgcgga agaagaggca acagttccaa acaataagat     120
cactgtagtg ggtgttggac aagttggtat ggcgtgtgct atcagcattc tgggaaagtc     180
tctggctgat gaacttgctc ttgtggatgt tttggaagat aagcttaaag gagaaatgat     240
ggatctgcag catgggagct atttcttca gacacctaaa attgtggcag ataaagatta      300
ttctgtgacc gccaattcta agattgtagt ggtaactgca ggagtccgtc agcaagaagg     360
ggagagtcgg ctcaatctgg tgcagagaaa tgttaatgtc ttcaaattca ttattcctca     420
gatcgtcaag tacagtcctg attgcatcat aattgtggtt tccaacccag tggacattct     480
tacgtatgtt acc                                                        493
```

    <210> 182
    <211> 209
    <212> PRT
    <213> homo sapien

    <400> 182
Ala Phe Ser Ser Asn Pro Lys Val Gln Val Glu Ala Ile Glu Gly Gly
-1   5   10   15
Ala Leu Gln Lys Leu Leu Val Ile Leu Ala Thr Glu Gln Pro Leu Thr
             20   25   30
Ala Lys Lys Lys Val Leu Phe Ala Leu Cys Ser Leu Leu Arg His Phe
         35   40   45
Pro Tyr Ala Gln Arg Gln Phe Leu Lys Leu Gly Gly Leu Gln Val Leu
     50   55   60
Arg Thr Leu Val Gln Glu Lys Gly Thr Glu Val Leu Ala Val Arg Val
65   70   75   80
Val Thr Leu Leu Tyr Asp Leu Val Thr Glu Lys Met Phe Ala Glu Glu
             85   90   95
Glu Ala Glu Leu Thr Gln Glu Met Ser Pro Glu Lys Leu Gln Gln Tyr
             100   105   110
Arg Gln Val His Leu Leu Pro Gly Leu Trp Glu Gln Gly Trp Cys Glu
         115   120   125
Ile Thr Ala His Leu Leu Ala Leu Pro Glu His Asp Ala Arg Glu Lys
     130   135   140
Val Leu Gln Thr Leu Gly Val Leu Leu Thr Thr Cys Arg Asp Arg Tyr
145   150   155   160
Arg Gln Asp Pro Gln Leu Gly Arg Thr Leu Ala Ser Leu Gln Ala Glu
             165   170   175
Tyr Gln Val Leu Ala Ser Leu Glu Leu Gln Asp Gly Glu Asp Glu Gly
             180   185   190
Tyr Phe Gln Glu Leu Leu Gly Ser Val Asn Ser Leu Leu Lys Glu Leu
         195   200   205
Arg

    <210> 183
    <211> 255
    <212> PRT
    <213> homo sapien

    <400> 183
Met Ala Ala Gly Val Glu Ala Ala Ala Glu Val Ala Ala Thr Glu Pro
1   5   10   15
```

```
Lys Met Glu Glu Glu Ser Gly Ala Pro Cys Val Pro Ser Gly Asn Gly
        20                  25                  30
Ala Pro Gly Pro Lys Gly Glu Glu Arg Pro Thr Gln Asn Glu Lys Arg
        35                  40                  45
Lys Glu Lys Asn Ile Lys Arg Gly Gly Asn Arg Phe Glu Pro Tyr Ser
    50                  55                  60
Asn Pro Thr Lys Arg Tyr Arg Ala Phe Ile Thr Asn Ile Pro Phe Asp
65                  70                  75                  80
Val Lys Trp Gln Ser Leu Lys Asp Leu Val Lys Glu Lys Val Gly Glu
                85                  90                  95
Val Thr Tyr Val Glu Leu Leu Met Asp Ala Glu Gly Lys Ser Arg Gly
            100                 105                 110
Cys Ala Val Val Glu Phe Lys Met Glu Glu Ser Met Lys Lys Ala Ala
        115                 120                 125
Glu Val Leu Asn Lys His Ser Leu Ser Gly Arg Pro Leu Lys Val Lys
    130                 135                 140
Glu Asp Pro Asp Gly Glu His Ala Arg Arg Ala Met Gln Lys Ala Gly
145                 150             --      155                 160
Arg Leu Gly Ser Thr Val Phe Val Ala Asn Leu Asp Tyr Lys Val Gly
            165                 170                 175
Trp Lys Lys Leu Lys Glu Val Phe Ser Met Ala Gly Val Val Val Arg
            180                 185                 190
Ala Asp Ile Leu Glu Asp Lys Asp Gly Lys Ser Arg Gly Ile Gly Ile
        195                 200                 205
Val Thr Phe Glu Gln Ser Ile Glu Ala Val Gln Ala Ile Ser Met Phe
    210                 215                 220
Asn Gly Gln Leu Leu Phe Asp Arg Pro Met His Val Lys Met Asp Glu
225                 230                 235                 240
Arg Ala Leu Pro Lys Gly Asp Phe Phe Pro Pro Glu Arg His Ser
                245                 250                 255
```

<210> 184
<211> 188
<212> PRT
<213> Homo sapien

<400> 184

```
Leu Ser Gly Ser Cys Ile Arg Arg Glu Gln Thr Pro Glu Lys Glu Lys
1               5                   10                  15
Gln Val Val Leu Phe Glu Glu Ala Ser Trp Thr Cys Thr Pro Ala Cys
        20                  25                  30
Gly Asp Glu Pro Arg Thr Val Ile Leu Leu Ser Ser Met Leu Ala Asp
        35                  40                  45
His Arg Leu Lys Leu Glu Asp Tyr Lys Asp Arg Leu Lys Ser Gly Glu
    50                  55                  60
His Leu Asn Pro Asp Gln Leu Glu Ala Val Glu Lys Tyr Glu Glu Val
65                  70                  75                  80
Leu His Asn Leu Glu Phe Ala Lys Glu Leu Gln Lys Thr Phe Ser Gly
                85                  90                  95
Leu Ser Leu Asp Leu Leu Lys Ala Gln Lys Lys Ala Gln Arg Arg Glu
            100                 105                 110
His Met Leu Lys Leu Glu Ala Glu Lys Lys Lys Leu Arg Thr Ile Leu
        115                 120                 125
Gln Val Gln Tyr Val Leu Gln Asn Leu Thr Gln Glu His Val Gln Lys
    130                 135                 140
```

```
Asp Phe Lys Gly Gly Leu Asn Gly Ala Val Tyr Leu Pro Ser Lys Glu
145                 150                 155                 160
Leu Asp Tyr Leu Ile Lys Phe Ser Lys Leu Thr Cys Pro Glu Arg Asn
                165                 170                 175
Glu Ser Leu Arg Gln Thr Leu Glu Gly Ser Thr Val
                180                 185
```

```
<210> 185
<211> 746
<212> PRT
<213> Homo sapien

<400> 185
```

```
Asp Lys His Leu Lys Asp Leu Leu Ser Lys Leu Leu Asn Ser Gly Tyr
1               5                   10                  15
Phe Glu Ser Ile Pro Val Pro Lys Asn Ala Lys Glu Lys Glu Val Pro
                20                  25                  30
Leu Glu Glu Glu Met Leu Ile Gln Ser Glu Lys Lys Thr Gln Leu Ser
        35                  40                  45
Lys Thr Glu Ser Val Lys Glu Ser Glu Ser Leu Met Glu Phe Ala Gln
        50                  55                  60
Pro Glu Ile Gln Pro Gln Glu Phe Leu Asn Arg Arg Tyr Met Thr Glu
65                  70                  75                  80
Val Asp Tyr Ser Asn Lys Gln Gly Glu Glu Gln Pro Trp Glu Ala Asp
                85                  90                  95
Tyr Ala Arg Lys Pro Asn Leu Pro Lys Arg Trp Asp Met Leu Thr Glu
            100                 105                 110
Pro Asp Gly Gln Glu Lys Lys Gln Glu Ser Phe Lys Ser Trp Glu Ala
        115                 120                 125
Ser Gly Lys His Gln Glu Val Ser Lys Pro Ala Val Ser Leu Glu Gln
    130                 135                 140
Arg Lys Gln Asp Thr Ser Lys Leu Arg Ser Thr Leu Pro Glu Glu Gln
145                 150                 155                 160
Lys Lys Gln Glu Ile Ser Lys Ser Lys Pro Ser Pro Ser Gln Trp Lys
                165                 170                 175
Gln Asp Thr Pro Lys Ser Lys Ala Gly Tyr Val Gln Glu Glu Gln Lys
            180                 185                 190
Lys Gln Glu Thr Pro Lys Leu Trp Pro Val Gln Leu Gln Lys Glu Gln
        195                 200                 205
Asp Pro Lys Lys Gln Thr Pro Lys Ser Trp Thr Pro Ser Met Gln Ser
    210                 215                 220
Glu Gln Asn Thr Thr Lys Ser Trp Thr Thr Pro Met Cys Glu Glu Gln
225                 230                 235                 240
Asp Ser Lys Gln Pro Glu Thr Pro Lys Ser Trp Glu Asn Asn Val Glu
                245                 250                 255
Ser Gln Lys His Ser Leu Thr Ser Gln Ser Gln Ile Ser Pro Lys Ser
            260                 265                 270
Trp Gly Val Ala Thr Ala Ser Leu Ile Pro Asn Asp Gln Leu Leu Pro
        275                 280                 285
Arg Lys Leu Asn Thr Glu Pro Lys Asp Val Pro Lys Pro Val His Gln
    290                 295                 300
Pro Val Gly Ser Ser Ser Thr Leu Pro Lys Asp Pro Val Leu Arg Lys
305                 310                 315                 320
Glu Lys Leu Gln Asp Leu Met Thr Gln Ile Gln Gly Thr Cys Asn Phe
                325                 330                 335
```

```
Met Gln Glu Ser Val Leu Asp Phe Asp Lys Pro Ser Ser Ala Ile Pro
            340                     345                 350
Thr Ser Gln Pro Pro Ser Ala Thr Pro Gly Ser Pro Val Ala Ser Lys
            355                 360                 365
Glu Gln Asn Leu Ser Ser Gln Ser Asp Phe Leu Gln Glu Pro Leu Gln
        370                 375                 380
Val Phe Asn Val Asn Ala Pro Leu Pro Pro Arg Lys Glu Gln Glu Ile
385                 390                 395                 400
Lys Glu Ser Pro Tyr Ser Pro Gly Tyr Asn Gln Ser Phe Thr Thr Ala
                405                 410                 415
Ser Thr Gln Thr Pro Pro Gln Cys Gln Leu Pro Ser Ile His Val Glu
            420                 425                 430
Gln Thr Val His Ser Gln Glu Thr Ala Ala Asn Tyr His Pro Asp Gly
        435                 440                 445
Thr Ile Gln Val Ser Asn Gly Ser Leu Ala Phe Tyr Pro Ala Gln Thr
    450                 455                 460
Asn Val Phe Pro Arg Pro Thr Gln Pro Phe Val Asn Ser Arg Gly Ser
465                 470                 475                 480
Val Arg Gly Cys Thr Arg Gly Gly Arg Leu Ile Thr Asn Ser Tyr Arg
                485                 490                 495
Ser Pro Gly Gly Tyr Lys Gly Phe Asp Thr Tyr Arg Gly Leu Pro Ser
            500                 505                 510
Ile Ser Asn Gly Asn Tyr Ser Gln Leu Gln Phe Gln Ala Arg Glu Tyr
        515                 520                 525
Ser Gly Ala Pro Tyr Ser Gln Arg Asp Asn Phe Gln Gln Cys Tyr Lys
    530                 535                 540
Arg Gly Gly Thr Ser Gly Gly Pro Arg Ala Asn Ser Arg Ala Gly Trp
545                 550                 555                 560
Ser Asp Ser Ser Gln Val Ser Ser Pro Glu Arg Asp Asn Glu Thr Phe
                565                 570                 575
Asn Ser Gly Asp Ser Gly Gln Gly Asp Ser Arg Ser Met Thr Pro Val
            580                 585                 590
Asp Val Pro Val Thr Asn Pro Ala Ala Thr Ile Leu Pro Val His Val
        595                 600                 605
Tyr Pro Leu Pro Gln Gln Met Arg Val Ala Phe Ser Ala Ala Arg Thr
    610                 615                 620
Ser Asn Leu Ala Pro Gly Thr Leu Asp Gln Pro Ile Val Phe Asp Leu
625                 630                 635                 640
Leu Leu Asn Asn Leu Gly Glu Thr Phe Asp Leu Gln Leu Gly Arg Phe
                645                 650                 655
Asn Cys Pro Val Asn Gly Thr Tyr Val Phe Ile Phe His Met Leu Lys
            660                 665                 670
Leu Ala Val Asn Val Pro Leu Tyr Val Asn Leu Met Lys Asn Glu Glu
        675                 680                 685
Val Leu Val Ser Ala Tyr Ala Asn Asp Gly Ala Pro Asp His Glu Thr
    690                 695                 700
Ala Ser Asn His Ala Ile Leu Gln Leu Phe Gln Gly Asp Gln Ile Trp
705                 710                 715                 720
Leu Arg Leu His Arg Gly Ala Ile Tyr Gly Ser Ser Trp Lys Tyr Ser
                725                 730                 735
Thr Phe Ser Gly Tyr Leu Leu Tyr Gln Asp
            740                 745
```

    <210> 186
    <211> 705

<212> PRT
<213> Homo sapien

<400> 186

```
Ala Leu Leu Asn Val Arg Gln Pro Pro Ser Thr Thr Thr Phe Val Leu
1               5                  10                 15
Asn Gln Ile Asn His Leu Pro Pro Leu Gly Ser Thr Ile Val Met Thr
            20                 25                 30
Lys Thr Pro Pro Val Thr Thr Asn Arg Gln Thr Ile Thr Leu Thr Lys
        35                 40                 45
Phe Ile Gln Thr Thr Ala Ser Thr Arg Pro Ser Val Ser Ala Pro Thr
    50                 55                 60
Val Arg Asn Ala Met Thr Ser Ala Pro Ser Lys Asp Gln Val Gln Leu
65                 70                 75                 80
Lys Asp Leu Leu Lys Asn Asn Ser Leu Asn Glu Leu Met Lys Leu Lys
                85                 90                 95
Pro Pro Ala Asn Ile Ala Gln Pro Val Ala Thr Ala Ala Thr Asp Val
            100                105                110
Ser Asn Gly Thr Val Lys Lys Glu Ser Ser Asn Lys Glu Gly Ala Arg
            115                120                125
Met Trp Ile Asn Asp Met Lys Met Arg Ser Phe Ser Pro Thr Met Lys
    130                135                140
Val Pro Val Val Lys Glu Asp Asp Glu Pro Glu Glu Glu Asp Glu Glu
145                150                155                160
Glu Met Gly His Ala Glu Thr Tyr Ala Glu Tyr Met Pro Ile Lys Leu
            165                170                175
Lys Ile Gly Leu Arg His Pro Asp Ala Val Val Glu Thr Ser Ser Leu
            180                185                190
Ser Ser Val Thr Pro Pro Asp Val Trp Tyr Lys Thr Ser Ile Ser Glu
        195                200                205
Glu Thr Ile Asp Asn Gly Trp Leu Ser Ala Leu Gln Leu Glu Ala Ile
    210                215                220
Thr Tyr Ala Ala Gln Gln His Glu Thr Phe Leu Pro Asn Gly Asp Arg
225                230                235                240
Ala Gly Phe Leu Ile Gly Asp Gly Ala Gly Val Gly Lys Gly Arg Thr
            245                250                255
Ile Ala Gly Ile Ile Tyr Glu Asn Tyr Leu Leu Ser Arg Lys Arg Ala
            260                265                270
Leu Trp Phe Ser Val Ser Asn Asp Leu Lys Tyr Asp Ala Glu Arg Asp
        275                280                285
Leu Arg Asp Ile Gly Ala Lys Asn Ile Leu Val His Ser Leu Asn Lys
    290                295                300
Phe Lys Tyr Gly Lys Ile Ser Ser Lys His Asn Gly Ser Val Lys Lys
305                310                315                320
Gly Val Ile Phe Ala Thr Tyr Ser Ser Leu Ile Gly Glu Ser Gln Ser
            325                330                335
Gly Gly Lys Tyr Lys Thr Arg Leu Lys Gln Leu Leu His Trp Cys Gly
        340                345                350
Asp Asp Phe Asp Gly Val Ile Val Phe Asp Glu Cys His Lys Ala Lys
        355                360                365
Asn Leu Cys Pro Val Gly Ser Ser Lys Pro Thr Lys Thr Gly Leu Ala
    370                375                380
Val Leu Glu Leu Gln Asn Lys Leu Pro Lys Ala Arg Val Val Tyr Ala
385                390                395                400
Ser Ala Thr Gly Ala Ser Glu Pro Arg Asn Met Ala Tyr Met Asn Arg
```

```
                    405                    410                    415
    Leu Gly Ile Trp Gly Glu Gly Thr Pro Phe Arg Glu Phe Ser Asp Phe
                420                    425                    430
    Ile Gln Ala Val Glu Arg Arg Gly Val Gly Ala Met Glu Ile Val Ala
                435                    440                    445
    Met Asp Met Lys Leu Arg Gly Met Tyr Ile Ala Arg Gln Leu Ser Phe
        450                    455                    460
    Thr Gly Val Thr Phe Lys Ile Glu Glu Val Leu Leu Ser Gln Ser Tyr
        465                    470                    475                    480
    Val Lys Met Tyr Asn Lys Ala Val Lys Leu Trp Val Ile Ala Arg Glu
                    485                    490                    495
    Arg Phe Gln Gln Ala Ala Asp Leu Ile Asp Ala Glu Gln Arg Met Lys
                500                    505                    510
    Lys Ser Met Trp Gly Gln Phe Trp Ser Ala His Gln Arg Phe Phe Lys
            515                    520                    525
    Tyr Leu Cys Ile Ala Ser Lys Val Lys Arg Val Val Gln Leu Ala Arg
        530                    535                    540
    Glu Glu Ile Lys Asn Gly Lys Cys Val Val Ile Gly Leu Gln Ser Thr
    545                    550                    555                    560
    Gly Glu Ala Arg Thr Leu Glu Ala Leu Glu Glu Gly Gly Gly Glu Leu
                565                    570                    575
    Asn Asp Phe Val Ser Thr Ala Lys Gly Val Leu Gln Ser Leu Ile Glu
                580                    585                    590
    Lys His Phe Pro Ala Pro Asp Arg Lys Lys Leu Tyr Ser Leu Leu Gly
        595                    600                    605
    Ile Asp Leu Thr Ala Pro Ser Asn Asn Ser Ser Pro Arg Asp Ser Pro
        610                    615                    620
    Cys Lys Glu Asn Lys Ile Lys Lys Arg Lys Gly Glu Glu Ile Thr Arg
    625                    630                    635                    640
    Glu Ala Lys Lys Ala Arg Lys Val Gly Gly Leu Thr Gly Ser Ser Ser
                645                    650                    655
    Asp Asp Ser Gly Ser Glu Ser Asp Ala Ser Asp Asn Glu Glu Ser Asp
                660                    665                    670
    Tyr Glu Ser Ser Lys Asn Met Ser Ser Gly Asp Asp Asp Asp Phe Asn
        675                    680                    685
    Pro Phe Leu Asp Glu Ser Asn Glu Asp Asp Glu Asn Asp Pro Trp Leu
        690                    695                    700
    Ile
    705


            <210> 187
            <211> 595
            <212> PRT
            <213> Homo sapien


        <400> 187
    Glu Ser Pro Arg His Arg Gly Glu Gly Gly Gly Glu Trp Gly Pro Gly
        1                    5                    10                    15
    Val Pro Arg Glu Arg Arg Glu Ser Ala Gly Glu Trp Gly Ala Asp Thr
                20                    25                    30
    Pro Lys Glu Gly Gly Glu Ser Ala Gly Glu Trp Gly Ala Glu Val Pro
            35                    40                    45
    Arg Gly Arg Gly Glu Gly Ala Gly Glu Trp Gly Pro Asp Thr Pro Lys
        50                    55                    60
    Glu Arg Gly Gln Gly Val Arg Glu Trp Gly Pro Glu Ile Pro Gln Glu
```

```
65                    70                   75              80
His Gly Glu Ala Thr Arg Asp Trp Ala Leu Glu Ser Pro Arg Ala Leu
                85                   90                   95
Gly Glu Asp Ala Arg Glu Leu Gly Ser Ser Pro His Asp Arg Gly Ala
             100                  105                  110
Ser Pro Arg Asp Leu Ser Gly Glu Ser Pro Cys Thr Gln Arg Ser Gly
             115                  120                  125
Leu Leu Pro Glu Arg Arg Gly Asp Ser Pro Trp Pro Pro Trp Pro Ser
        130                  135                  140
Pro Gln Glu Arg Asp Ala Gly Thr Arg Asp Arg Glu Glu Ser Pro Arg
145                  150                  155                  160
Asp Trp Gly Gly Ala Glu Ser Pro Arg Gly Trp Glu Ala Gly Pro Arg
                165                  170                  175
Glu Trp Gly Pro Ser Pro Ser Gly His Gly Asp Gly Pro Arg Arg Arg
             180                  185                  190
Pro Arg Lys Arg Arg Gly Arg Lys Gly Arg Met Gly Arg Gln His Glu
        195                  200                  205
Ala Ala Ala Thr Ala Ala Thr Ala Ala Thr Ala Thr Gly Gly Thr Ala
    210                  215                  220
Glu Glu Ala Gly Ala Ser Ala Pro Glu Ser Gln Ala Gly Gly Gly Pro
225                  230                  235                  240
Arg Gly Arg Ala Arg Gly Pro Arg Gln Gln Gly Arg Arg Arg His Gly
             245                  250                  255
Thr Gln Arg Arg Arg Gly Pro Pro Gln Ala Arg Glu Glu Gly Pro Arg
             260                  265                  270
Asp Ala Thr Thr Ile Leu Gly Leu Gly Thr Pro Ser Gly Glu Gln Arg
             275                  280                  285
Ala Asp Gln Ser Gln Ala Leu Pro Ala Leu Ala Gly Ala Ala Ala Ala
    290                  295                  300
His Ala His Ala Ile Pro Gly Ala Gly Pro Ala Ala Ala Pro Val Gly
305                  310                  315                  320
Gly Arg Gly Arg Arg Gly Gly Trp Arg Gly Gly Arg Arg Gly Gly Ser
             325                  330                  335
Ala Gly Ala Gly Gly Gly Gly Arg Gly Gly Arg Gly Arg Gly Arg Gly
        340                  345                  350
Gly Gly Arg Gly Gly Gly Gly Ala Gly Arg Gly Gly Gly Ala Ala Gly
        355                  360                  365
Pro Arg Glu Gly Ala Ser Ser Pro Gly Ala Arg Arg Gly Glu Gln Arg
    370                  375                  380
Arg Arg Gly Arg Gly Pro Pro Ala Ala Gly Ala Ala Gln Val Ser Ala
385                  390                  395                  400
Arg Gly Arg Arg Ala Arg Gly Gln Arg Ala Gly Glu Glu Ala Gln Asp
             405                  410                  415
Gly Leu Leu Pro Arg Gly Arg Asp Arg Leu Pro Leu Arg Pro Gly Asp
        420                  425                  430
Ala Asn Gln Arg Ala Glu Arg Pro Gly Pro Pro Arg Gly Gly His Gly
    435                  440                  445
Pro Val Asn Ala Ser Ser Ala Pro Asp Thr Ser Pro Pro Arg His Pro
    450                  455                  460
Arg Arg Trp Val Ser Gln Gln Arg Gln Arg Leu Trp Arg Gln Phe Arg
465                  470                  475                  480
Val Gly Gly Gly Phe Pro Pro Pro Pro Ser Arg Pro Pro Ala Val
             485                  490                  495
Leu Leu Pro Leu Leu Arg Leu Ala Cys Ala Gly Asp Pro Gly Ala Thr
        500                  505                  510
```

```
Arg Pro Gly Pro Arg Arg Pro Ala Arg Arg Pro Arg Gly Glu Leu Ile
        515             520             525
Pro Arg Arg Pro Asp Pro Ala Ala Pro Ser Glu Glu Gly Leu Arg Met
        530             535             540
Glu Ser Ser Val Asp Asp Gly Ala Thr Ala Thr Thr Ala Asp Ala Ala
545             550             555             560
Ser Gly Glu Ala Pro Glu Ala Gly Pro Ser Pro Ser His Ser Pro Thr
                565             570             575
Met Cys Gln Thr Gly Gly Pro Gly Pro Pro Pro Gln Pro Pro Arg
        580             585             590
Trp Leu Pro
        595


        <210> 188
        <211> 376
        <212> PRT
        <213> Homo sapien


        <400> 188
Glu Met Arg Lys Phe Asp Val Pro Ser Met Glu Ser Thr Leu Asn Gln
1               5               10              15
Pro Ala Met Leu Glu Thr Leu Tyr Ser Asp Pro His Tyr Arg Ala His
            20              25              30
Phe Pro Asn Pro Arg Pro Asp Thr Asn Lys Asp Val Tyr Lys Val Leu
        35              40                  45
Pro Glu Ser Lys Lys Ala Pro Gly Ser Gly Ala Val Phe Glu Arg Asn
    50              55                  60
Gly Pro His Ala Ser Ser Ser Gly Val Leu Pro Leu Gly Leu Gln Pro
65                  70                  75                  80
Ala Pro Gly Leu Ser Lys Ser Leu Ser Ser Gln Val Trp Gln Pro Ser
                85                  90                  95
Pro Asp Pro Trp His Pro Gly Glu Gln Ser Cys Glu Leu Ser Thr Cys
            100             105             110
Arg Gln Gln Leu Glu Leu Ile Arg Leu Gln Met Glu Gln Met Gln Leu
        115             120             125
Gln Asn Gly Ala Met Cys His His Pro Ala Ala Phe Ala Pro Leu Leu
        130             135             140
Pro Thr Leu Glu Pro Ala Gln Trp Leu Ser Ile Leu Asn Ser Asn Glu
145             150             155             160
His Leu Leu Lys Glu Lys Glu Leu Leu Ile Asp Lys Gln Arg Lys His
                165             170             175
Ile Ser Gln Leu Glu Gln Lys Val Arg Glu Ser Glu Leu Gln Val His
            180             185             190
Ser Ala Leu Leu Gly Arg Pro Ala Pro Phe Gly Asp Val Cys Leu Leu
        195             200             205
Arg Leu Gln Glu Leu Gln Arg Glu Asn Thr Phe Leu Arg Ala Gln Phe
        210             215             220
Ala Gln Lys Thr Glu Ala Leu Ser Lys Glu Lys Met Glu Leu Glu Lys
225             230             235             240
Lys Leu Ser Ala Ser Glu Val Glu Ile Gln Leu Ile Arg Glu Ser Leu
                245             250             255
Lys Val Thr Leu Gln Lys His Ser Glu Glu Gly Lys Lys Gln Glu Glu
        260             265             270
Arg Val Lys Gly Arg Asp Lys His Ile Asn Asn Leu Lys Lys Lys Cys
        275             280             285
```

```
Gln Lys Glu Ser Glu Gln Asn Arg Glu Lys Gln Gln Arg Ile Glu Thr
    290                 295                 300
Leu Glu Arg Tyr Leu Ala Asp Leu Pro Thr Leu Glu Asp His Gln Lys
305                 310                 315                 320
Gln Thr Glu Gln Leu Lys Asp Ala Glu Leu Lys Asn Thr Glu Leu Gln
                325                 330                 335
Glu Arg Val Ala Glu Leu Glu Thr Leu Leu Glu Asp Thr Gln Ala Thr
            340                 345                 350
Cys Arg Glu Lys Glu Val Gln Leu Glu Ser Leu Arg Gln Arg Glu Ala
        355                 360                 365
Asp Leu Ser Ser Ala Arg His Arg
370                 375


        <210> 189
        <211> 160
        <212> PRT
        <213> Homo sapien


        <400> 189
Met Leu Glu Ala His Arg Arg Gln Arg His Pro Phe Leu Leu Leu Gly
1                   5                   10                  15
Thr Thr Ala Asn Arg Thr Gln Ser Leu Asn Tyr Gly Cys Ile Val Glu
            20                  25                  30
Asn Pro Gln Thr His Glu Val Leu His Tyr Val Glu Lys Pro Ser Thr
        35                  40                  45
Phe Ile Ser Asp Ile Ile Asn Cys Gly Ile Tyr Leu Phe Ser Pro Glu
        50                  55                  60
Ala Leu Lys Pro Leu Arg Asp Val Phe Gln Arg Asn Gln Gln Asp Gly
65                  70                  75                  80
Gln Leu Glu Asp Ser Pro Gly Leu Trp Pro Gly Ala Gly Thr Ile Arg
                85                  90                  95
Leu Glu Gln Asp Val Phe Ser Ala Leu Ala Gly Gln Gly Gln Ile Tyr
                100                 105                 110
Val His Leu Thr Asp Gly Ile Trp Ser Gln Ile Lys Ser Ala Gly Ser
        115                 120                 125
Ala Leu Tyr Ala Ser Arg Leu Tyr Leu Ser Arg Tyr Gln Asp Thr His
    130                 135                 140
Pro Glu Arg Leu Ala Lys His Thr Pro Gly Gly Pro Trp Ile Arg Gly
145                 150                 155                 160


        <210> 190
        <211> 146
        <212> PRT
        <213> Homo sapien


        <400> 190
Met Asp Pro Arg Ala Ser Leu Leu Leu Leu Gly Asn Val Tyr Ile His
1                   5                   10                  15
Pro Thr Ala Lys Val Ala Pro Ser Ala Val Leu Gly Pro Asn Val Ser
            20                  25                  30
Ile Gly Lys Gly Val Thr Val Gly Glu Gly Val Arg Leu Arg Glu Ser
        35                  40                  45
Ile Val Leu His Gly Ala Thr Leu Gln Glu His Thr Cys Val Leu His
    50                  55                  60
Ser Ile Val Gly Trp Gly Ser Thr Val Gly Arg Trp Ala Arg Val Glu
```

```
      65                    70                    75                    80
    Gly Thr Pro Ser Asp Pro Asn Pro Asn Asp Pro Arg Ala Arg Met Asp
                          85                    90                    95
    Ser Glu Ser Leu Phe Lys Asp Gly Lys Leu Leu Pro Ala Ile Thr Ile
                          100                   105                   110
    Leu Gly Cys Arg Val Arg Ile Pro Ala Glu Val Leu Ile Leu Asn Ser
                          115                   120                   125
    Ile Val Leu Pro His Lys Glu Leu Ser Arg Ser Phe Thr Asn Gln Ile
        130                   135                   140
    Ile Leu
    145


        <210> 191
        <211> 704
        <212> PRT
        <213> Homo sapien


        <400> 191
    Glu Gly Gly Cys Ala Ala Gly Arg Gly Arg Glu Leu Glu Pro Glu Leu
    1                   5                   10                    15
    Glu Pro Gly Pro Gly Pro Gly Ser Ala Leu Glu Pro Gly Glu Glu Phe
                          20                    25                    30
    Glu Ile Val Asp Arg Ser Gln Leu Pro Gly Pro Gly Asp Leu Arg Ser
        35                    40                    45
    Ala Thr Arg Pro Arg Ala Ala Glu Gly Trp Ser Ala Pro Ile Leu Thr
        50                    55                    60
    Leu Ala Arg Arg Ala Thr Gly Asn Leu Ser Ala Ser Cys Gly Ser Ala
    65                    70                    75                    80
    Leu Arg Ala Ala Ala Gly Leu Gly Gly Gly Asp Ser Gly Asp Gly Thr
                          85                    90                    95
    Ala Arg Ala Ala Ser Lys Cys Gln Met Met Glu Glu Arg Ala Asn Leu
                  100                   105                   110
    Met His Met Met Lys Leu Ser Ile Lys Val Leu Leu Gln Ser Ala Leu
        115                   120                   125
    Ser Leu Gly Arg Ser Leu Asp Ala Asp His Ala Pro Leu Gln Gln Phe
        130                   135                   140
    Phe Val Val Met Glu His Cys Leu Lys His Gly Leu Lys Val Lys Lys
    145                   150                   155                   160
    Ser Phe Ile Gly Gln Asn Lys Ser Phe Phe Gly Pro Leu Glu Leu Val
                          165                   170                   175
    Glu Lys Leu Cys Pro Glu Ala Ser Asp Ile Ala Thr Ser Val Arg Asn
                  180                   185                   190
    Leu Pro Glu Leu Lys Thr Ala Val Gly Arg Gly Arg Ala Trp Leu Tyr
        195                   200                   205
    Leu Ala Leu Met Gln Lys Lys Leu Ala Asp Tyr Leu Lys Val Leu Ile
        210                   215                   220
    Asp Asn Lys His Leu Leu Ser Glu Phe Tyr Glu Pro Glu Ala Leu Met
    225                   230                   235                   240
    Met Glu Glu Glu Gly Met Val Ile Val Gly Leu Leu Val Gly Leu Asn
                          245                   250                   255
    Val Leu Asp Ala Asn Leu Cys Leu Lys Gly Glu Asp Leu Asp Ser Gln
                  260                   265                   270
    Val Gly Val Ile Asp Phe Ser Leu Tyr Leu Lys Asp Val Gln Asp Leu
        275                   280                   285
    Asp Gly Gly Lys Glu His Glu Arg Ile Thr Asp Val Leu Asp Gln Lys
```

```
        290                    295                    300
Asn Tyr Val Glu Glu Leu Asn Arg His Leu Ser Cys Thr Val Gly Asp
305                    310                    315                    320
Leu Gln Thr Lys Ile Asp Gly Leu Glu Lys Thr Asn Ser Lys Leu Gln
                325                    330                    335
Glu Glu Leu Ser Ala Ala Thr Asp Arg Ile Cys Ser Leu Gln Glu Glu
            340                    345                    350
Gln Gln Gln Leu Arg Glu Gln Asn Glu Leu Ile Arg Glu Arg Ser Glu
        355                    360                    365
Lys Ser Val Glu Ile Thr Lys Gln Asp Thr Lys Val Glu Leu Glu Thr
    370                    375                    380
Tyr Lys Gln Thr Arg Gln Gly Leu Asp Glu Met Tyr Ser Asp Val Trp
385                    390                    395                    400
Lys Gln Leu Lys Glu Glu Lys Lys Val Arg Leu Glu Leu Glu Lys Glu
                405                    410                    415
Leu Glu Leu Gln Ile Gly Met Lys Thr Glu Met Glu Ile Ala Met Lys
            420                    425                    430
Leu Leu Glu Lys Asp Thr His Glu Lys Gln Asp Thr Leu Val Ala Leu
        435                    440                    445
Arg Gln Gln Leu Glu Glu Val Lys Ala Ile Asn Leu Gln Met Phe His
    450                    455                    460
Lys Ala Gln Asn Ala Glu Ser Ser Leu Gln Gln Lys Asn Glu Ala Ile
465                    470                    475                    480
Thr Ser Phe Glu Gly Lys Thr Asn Gln Val Met Ser Ser Met Lys Gln
                485                    490                    495
Met Glu Glu Arg Leu Gln His Ser Glu Arg Ala Arg Gln Gly Ala Glu
            500                    505                    510
Glu Arg Ser His Lys Leu Gln Gln Glu Leu Gly Gly Arg Ile Gly Ala
        515                    520                    525
Leu Gln Leu Gln Leu Ser Gln Leu His Glu Gln Cys Ser Ser Leu Glu
    530                    535                    540
Lys Glu Leu Lys Ser Glu Lys Glu Gln Arg Gln Ala Leu Gln Arg Glu
545                    550                    555                    560
Leu Gln His Glu Lys Asp Thr Ser Ser Leu Leu Arg Met Glu Leu Gln
                565                    570                    575
Gln Val Glu Gly Leu Lys Lys Glu Leu Arg Glu Leu Gln Asp Glu Lys
            580                    585                    590
Ala Glu Leu Gln Lys Ile Cys Glu Glu Gln Glu Gln Ala Leu Gln Glu
        595                    600                    605
Met Gly Leu His Leu Ser Gln Ser Lys Leu Lys Met Glu Asp Ile Lys
    610                    615                    620
Glu Val Asn Gln Ala Leu Lys Gly His Ala Trp Leu Lys Asp Asp Glu
625                    630                    635                    640
Ala Thr His Cys Arg Gln Cys Glu Lys Glu Phe Ser Ile Ser Arg Arg
                645                    650                    655
Lys His His Cys Arg Asn Cys Gly His Ile Phe Cys Asn Thr Cys Ser
            660                    665                    670
Ser Asn Glu Leu Ala Leu Pro Ser Tyr Pro Lys Pro Val Arg Val Cys
        675                    680                    685
Asp Ser Cys His Thr Leu Leu Leu Gln Arg Cys Ser Ser Thr Ala Ser
    690                    695                    700
```

<210> 192
<211> 331
<212> PRT

100

<213> Homo sapien

<400> 192
Arg Ala Gly Ala Ser Ala Met Ala Leu Arg Lys Glu Leu Leu Lys Ser
1   5   10   15
Ile Trp Tyr Ala Phe Thr Ala Leu Asp Val Glu Lys Ser Gly Lys Val
            20   25   30
Ser Lys Ser Gln Leu Lys Val Leu Ser His Asn Leu Tyr Thr Val Leu
        35   40   45
His Ile Pro His Asp Pro Val Ala Leu Glu Glu His Phe Arg Asp Asp
    50   55   60
Asp Asp Gly Pro Val Ser Ser Gln Gly Tyr Met Pro Tyr Leu Asn Lys
65   70   75   80
Tyr Ile Leu Asp Lys Val Glu Glu Gly Ala Phe Val Lys Glu His Phe
                85   90   95
Asp Glu Leu Cys Trp Thr Leu Thr Ala Lys Lys Asn Tyr Arg Ala Asp
            100   105   110
Ser Asn Gly Asn Ser Met Leu Ser Asn Gln Asp Ala Phe Arg Leu Trp
        115   120   125
Cys Leu Phe Asn Phe Leu Ser Glu Asp Lys Tyr Pro Leu Ile Met Val
    130   135   140
Pro Asp Glu Val Glu Tyr Leu Leu Lys Lys Val Leu Ser Ser Met Ser
145   150   155   160
Leu Glu Val Ser Leu Gly Glu Leu Glu Glu Leu Leu Ala Gln Glu Ala
                165   170   175
Gln Val Ala Gln Thr Thr Gly Gly Leu Ser Val Trp Gln Phe Leu Glu
            180   185   190
Leu Phe Asn Ser Gly Arg Cys Leu Arg Gly Val Gly Arg Asp Thr Leu
        195   200   205
Ser Met Ala Ile His Glu Val Tyr Gln Glu Leu Ile Gln Asp Val Leu
    210   215   220
Lys Gln Gly Tyr Leu Trp Lys Arg Gly His Leu Arg Arg Asn Trp Ala
225   230   235   240
Glu Arg Trp Phe Gln Leu Gln Pro Ser Cys Leu Cys Tyr Phe Gly Ser
                245   250   255
Glu Glu Cys Lys Glu Lys Arg Gly Ile Ile Pro Leu Asp Ala His Cys
            260   265   270
Cys Val Glu Val Leu Pro Asp Arg Asp Gly Lys Arg Cys Met Phe Cys
        275   280   285
Val Lys Thr Ala Thr Arg Thr Tyr Glu Met Ser Ala Ser Asp Thr Arg
    290   295   300
Gln Arg Gln Glu Trp Thr Ala Ala Ile Gln Met Ala Ile Arg Leu Gln
305   310   315   320
Ala Glu Gly Lys Thr Ser Leu His Lys Asp Leu
                325   330

<210> 193
<211> 475
<212> PRT
<213> Homo sapien

<400> 193
Lys Asn Ser Pro Leu Leu Ser Val Ser Ser Gln Thr Ile Thr Lys Glu
1   5   10   15
Asn Asn Arg Asn Val His Leu Glu His Ser Glu Gln Asn Pro Gly Ser

```
                 20                    25                    30
Ser Ala Gly Asp Thr Ser Ala Ala His Gln Val Val Leu Gly Glu Asn
         35                    40                    45
Leu Ile Ala Thr Ala Leu Cys Leu Ser Gly Ser Gly Ser Gln Ser Asp
         50                    55                    60
Leu Lys Asp Val Ala Ser Thr Ala Gly Glu Glu Gly Asp Thr Ser Leu
65                    70                    75                    80
Arg Glu Ser Leu His Pro Val Thr Arg Ser Leu Lys Ala Gly Cys His
                 85                    90                    95
Thr Lys Gln Leu Ala Ser Arg Asn Cys Ser Glu Glu Lys Ser Pro Gln
                 100                   105                   110
Thr Ser Ile Leu Lys Glu Gly Asn Arg Asp Thr Ser Leu Asp Phe Arg
         115                   120                   125
Pro Val Val Ser Pro Ala Asn Gly Val Glu Gly Val Arg Val Asp Gln
    130                   135                   140
Asp Asp Asp Gln Asp Ser Ser Ser Leu Lys Leu Ser Gln Asn Ile Ala
145                   150                   155                   160
Val Gln Thr Asp Phe Lys Thr Ala Asp Ser Glu Val Asn Thr Asp Gln
                 165                   170                   175
Asp Ile Glu Lys Asn Leu Asp Lys Met Met Thr Glu Arg Thr Leu Leu
                 180                   185                   190
Lys Glu Arg Tyr Gln Glu Val Leu Asp Lys Gln Arg Gln Val Glu Asn
         195                   200                   205
Gln Leu Gln Val Gln Leu Lys Gln Leu Gln Gln Arg Arg Glu Glu Glu
    210                   215                   220
Met Lys Asn His Gln Glu Ile Leu Lys Ala Ile Gln Asp Val Thr Ile
225                   230                   235                   240
Lys Arg Glu Glu Thr Lys Lys Lys Ile Glu Lys Glu Lys Lys Glu Phe
                 245                   250                   255
Leu Gln Lys Glu Gln Asp Leu Lys Ala Glu Ile Glu Lys Leu Cys Glu
                 260                   265                   270
Lys Gly Arg Arg Glu Val Trp Glu Met Glu Leu Asp Arg Leu Lys Asn
         275                   280                   285
Gln Asp Gly Glu Ile Asn Arg Asn Ile Met Glu Glu Thr Glu Arg Ala
    290                   295                   300
Trp Lys Ala Glu Ile Leu Ser Leu Glu Ser Arg Lys Glu Leu Leu Val
305                   310                   315                   320
Leu Lys Leu Glu Glu Ala Glu Lys Glu Ala Glu Leu His Leu Thr Tyr
                 325                   330                   335
Leu Lys Ser Thr Pro Pro Thr Leu Glu Thr Val Arg Ser Lys Gln Glu
         340                   345                   350
Trp Glu Thr Arg Leu Asn Gly Val Arg Ile Met Lys Lys Asn Val Arg
         355                   360                   365
Asp Gln Phe Asn Ser His Ile Gln Leu Val Arg Asn Gly Ala Lys Leu
    370                   375                   380
Ser Ser Leu Pro Gln Ile Pro Thr Pro Thr Leu Pro Pro Pro Pro Ser
385                   390                   395                   400
Glu Thr Asp Phe Met Leu Gln Val Phe Gln Pro Ser Pro Ser Leu Ala
                 405                   410                   415
Pro Arg Met Pro Phe Ser Ile Gly Gln Val Thr Met Pro Met Val Met
                 420                   425                   430
Pro Ser Ala Asp Pro Arg Ser Leu Ser Phe Pro Ile Leu Asn Pro Ala
         435                   440                   445
Leu Ser Gln Pro Ser Gln Pro Ser Ser Pro Leu Pro Gly Ser His Gly
    450                   455                   460
```

Arg Asn Ser Pro Gly Leu Gly Ser Leu Val Ser
465   470   475

<210> 194
<211> 241
<212> PRT
<213> Homo sapien

<400> 194
Met Ser Gly Glu Ser Ala Arg Ser Leu Gly Lys Gly Ser Ala Pro Pro
1   5   10   15
Gly Pro Val Pro Glu Gly Ser Ile Arg Ile Tyr Ser Met Arg Phe Cys
            20   25   30
Pro Phe Ala Glu Arg Thr Arg Leu Val Leu Lys Ala Lys Gly Ile Arg
        35   40   45
His Glu Val Ile Asn Ile Asn Leu Lys Asn Lys Pro Glu Trp Phe Phe
    50   55   60
Lys Lys Asn Pro Phe Gly Leu Val Pro Val Leu Glu Asn Ser Gln Gly
65   70   75   80
Gln Leu Ile Tyr Glu Ser Ala Ile Thr Cys Glu Tyr Leu Asp Glu Ala
                85   90   95
Tyr Pro Gly Lys Lys Leu Leu Pro Asp Asp Pro Tyr Glu Lys Ala Cys
                100   105   110
Gln Lys Met Ile Leu Glu Leu Phe Ser Lys Val Pro Ser Leu Val Gly
        115   120   125
Ser Phe Ile Arg Ser Gln Asn Lys Glu Asp Tyr Ala Gly Leu Lys Glu
    130   135   140
Glu Phe Arg Lys Glu Phe Thr Lys Leu Glu Glu Val Leu Thr Asn Lys
145   150   155   160
Lys Thr Thr Phe Phe Gly Gly Asn Ser Ile Ser Met Ile Asp Tyr Leu
                165   170   175
Ile Trp Pro Trp Phe Glu Arg Leu Glu Ala Met Lys Leu Asn Glu Cys
            180   185   190
Val Asp His Thr Pro Lys Leu Lys Leu Trp Met Ala Ala Met Lys Glu
        195   200   205
Asp Pro Thr Val Ser Ala Leu Leu Thr Ser Glu Lys Asp Trp Gln Gly
    210   215   220
Phe Leu Glu Leu Tyr Leu Gln Asn Ser Pro Glu Ala Cys Asp Tyr Gly
225   230   235   240
Leu

<210> 195
<211> 138
<212> PRT
<213> Homo sapien

<400> 195
Gln Thr Lys Ile Leu Glu Glu Asp Leu Glu Gln Ile Lys Leu Ser Leu
1   5   10   15
Arg Glu Arg Gly Arg Glu Leu Thr Thr Gln Arg Gln Leu Met Gln Glu
            20   25   30
Arg Ala Glu Glu Gly Lys Gly Pro Ser Lys Ala Gln Arg Gly Ser Leu
        35   40   45
Glu His Met Lys Leu Ile Leu Arg Asp Lys Glu Lys Glu Val Glu Cys

```
        50                  55                    60
Gln Gln Glu His Ile His Glu Leu Gln Glu Leu Lys Asp Gln Leu Glu
65                  70                    75                    80
Gln Gln Leu Gln Gly Leu His Arg Lys Val Gly Glu Thr Ser Leu Leu
                85                    90                    95
Leu Ser Gln Arg Glu Gln Glu Ile Val Val Leu Gln Gln Gln Leu Gln
                100                   105                   110
Glu Ala Arg Glu Gln Gly Glu Leu Lys Glu Gln Ser Leu Gln Ser Gln
        115                   120                   125
Leu Asp Glu Ala Gln Arg Ala Leu Ala Gln
        130                   135
```

```
        <210> 196
        <211> 102
        <212> PRT
        <213> Homo sapien


        <400> 196
Met Ser Lys Arg Lys Ala Pro Gln Glu Thr Leu Asn Gly Gly Ile Thr
1                   5                     10                    15
Asp Met Leu Thr Glu Leu Ala Asn Phe Glu Lys Asn Val Ser Gln Ala
                20                    25                    30
Ile His Lys Tyr Asn Ala Tyr Arg Lys Ala Ala Ser Val Ile Ala Lys
        35                    40                    45
Tyr Pro His Lys Ile Lys Ser Gly Ala Glu Ala Lys Lys Leu Pro Gly
        50                    55                    60
Val Gly Thr Lys Ile Ala Glu Lys Ile Asp Glu Phe Leu Ala Thr Gly
65                  70                    75                    80
Lys Leu Arg Lys Leu Glu Lys Ile Arg Gln Asp Asp Thr Ser Ser Ser
                85                    90                    95
Ile Asn Phe Leu Thr Arg
                100
```

```
        <210> 197
        <211> 138
        <212> PRT
        <213> Homo sapien


        <400> 197
Glu Ala Asn Glu Val Thr Asp Ser Ala Tyr Met Gly Ser Glu Ser Thr
1                   5                     10                    15
Tyr Ser Glu Cys Glu Thr Phe Thr Asp Glu Asp Thr Ser Thr Leu Val
                20                    25                    30
His Pro Glu Leu Gln Pro Glu Gly Asp Ala Asp Ser Ala Gly Gly Ser
        35                    40                    45
Ala Val Pro Ser Glu Cys Leu Asp Ala Met Glu Glu Pro Asp His Gly
        50                    55                    60
Ala Leu Leu Leu Leu Pro Gly Arg Pro His Pro His Gly Gln Ser Val
65                  70                    75                    80
Ile Thr Val Ile Gly Gly Glu Glu His Phe Glu Asp Tyr Gly Glu Gly
                85                    90                    95
Ser Glu Ala Glu Leu Ser Pro Glu Thr Leu Cys Asn Gly Gln Leu Gly
                100                   105                   110
Cys Ser Asp Pro Ala Phe Leu Thr Pro Ser Pro Thr Lys Arg Leu Ser
        115                   120                   125
```

Ser Lys Lys Val Ala Arg Tyr Leu His Gln
    130   135

        <210> 198
        <211> 100
        <212> PRT
        <213> Homo sapien

        <400> 198
Met Gly Asp Val Lys Asn Phe Leu Tyr Ala Trp Cys Gly Lys Arg Lys
1   5   10   15
Met Thr Pro Ser Tyr Glu Ile Arg Ala Val Gly Asn Lys Asn Arg Gln
                20   25   30
Lys Phe Met Cys Glu Val Gln Val Glu Gly Tyr Asn Tyr Thr Gly Met
        35   40   45
Gly Asn Ser Thr Asn Lys Lys Asp Ala Gln Ser Asn Ala Ala Arg Asp
    50   55   60
Phe Val Asn Tyr Leu Val Arg Ile Asn Glu Ile Lys Ser Glu Glu Val
65   70   75   80
Pro Ala Phe Gly Val Ala Ser Pro Pro Pro Leu Thr Asp Thr Pro Asp
                85   90   95
Thr Thr Ala Asn
            100

        <210> 199
        <211> 127
        <212> PRT
        <213> Homo sapien

        <400> 199
Met Val Lys Glu Thr Thr Tyr Tyr Asp Val Leu Gly Val Lys Pro Asn
1   5   10   15
Ala Thr Gln Glu Glu Leu Lys Lys Ala Tyr Arg Lys Leu Ala Leu Lys
                20   25   30
Tyr His Pro Asp Lys Asn Pro Asn Glu Gly Glu Lys Phe Lys Gln Ile
        35   40   45
Ser Gln Ala Tyr Glu Val Leu Ser Asp Ala Lys Lys Arg Glu Leu Tyr
    50   55   60
Asp Lys Gly Gly Glu Gln Ala Ile Lys Glu Gly Gly Ala Gly Gly Gly
65   70   75   80
Phe Gly Ser Pro Met Asp Ile Phe Asp Met Phe Phe Gly Gly Gly Gly
                85   90   95
Arg Met Gln Arg Glu Arg Arg Gly Lys Asn Val Val His Gln Leu Ser
            100   105   110
Val Thr Leu Glu Asp Leu Tyr Asn Gly Ala Thr Arg Lys Leu Ala
        115   120   125

        <210> 200
        <211> 90
        <212> PRT
        <213> Homo sapien

        <400> 200
Met Ala Cys Pro Leu Asp Gln Ala Ile Gly Leu Leu Val Ala Ile Phe
1   5   10   15

```
His Lys Tyr Ser Gly Arg Glu Gly Asp Lys His Thr Leu Ser Lys Lys
        20                      25                  30
Glu Leu Lys Glu Leu Ile Gln Lys Glu Leu Thr Ile Gly Ser Lys Leu
        35                      40                  45
Gln Asp Ala Glu Ile Ala Arg Leu Met Glu Asp Leu Asp Arg Asn Lys
    50                      55                  60
Asp Gln Glu Val Asn Phe Gln Glu Tyr Val Thr Phe Leu Gly Ala Leu
65                      70                  75                  80
Ala Leu Ile Tyr Asn Glu Ala Leu Lys Gly
                85                      90
```

        <210> 201
        <211> 120
        <212> PRT
        <213> Homo sapien

        <400> 201
```
Met Glu Thr Pro Ser Gln Arg Arg Ala Thr Arg Ser Gly Ala Gln Ala
.1              5                   10                  15
Ser Ser Thr Pro Leu Ser Pro Thr Arg Ile Thr Arg Leu Gln Glu Lys
            20                  25                  30
Glu Asp Leu Gln Glu Leu Asn Asp Arg Leu Ala Val Tyr Ile Asp Arg
        35                  40                  45
Val Arg Ser Leu Glu Thr Glu Asn Ala Gly Leu Arg Leu Arg Ile Thr
    50                  55                  60
Glu Ser Glu Glu Val Val Ser Arg Glu Val Ser Gly Ile Lys Ala Ala
65                  70                  75                  80
Tyr Glu Ala Glu Leu Gly Asp Ala Arg Lys Thr Leu Asp Ser Val Ala
                85                  90                  95
Lys Glu Arg Ala Arg Leu Gln Leu Glu Leu Ser Lys Val Arg Glu Glu
            100                 105                 110
Phe Lys Glu Leu Lys Ala Arg Asn
        115                 120
```

        <210> 202
        <211> 177
        <212> PRT
        <213> Homo sapien

        <400> 202
```
Met Ala Ala Gly Val Glu Ala Ala Glu Val Ala Ala Thr Glu Ile
1               5                   10                  15
Lys Met Glu Glu Glu Ser Gly Ala Pro Gly Val Pro Ser Gly Asn Gly
        20                  25                  30
Ala Pro Gly Pro Lys Gly Glu Gly Glu Arg Pro Ala Gln Asn Glu Lys
        35                  40                  45
Arg Lys Glu Lys Asn Ile Lys Arg Gly Gly Asn Arg Phe Glu Pro Tyr
    50                  55                  60
Ala Asn Pro Thr Lys Arg Tyr Arg Ala Phe Ile Thr Asn Ile Pro Phe
65                  70                  75                  80
Asp Val Lys Trp Gln Ser Leu Lys Asp Leu Val Lys Glu Lys Val Gly
                85                  90                  95
Glu Val Thr Tyr Val Glu Leu Leu Met Asp Ala Glu Gly Lys Ser Arg
            100                 105                 110
Gly Cys Ala Val Val Glu Phe Lys Met Glu Glu Ser Met Lys Lys Ala
```

```
                115                   120                   125
Ala Glu Val Leu Asn Lys His Ser Leu Ser Gly Arg Pro Leu Lys Val
    130                   135                   140
Lys Glu Asp Pro Asp Gly Glu His Ala Arg Arg Ala Met Gln Lys Ala
145                   150                   155                   160
Gly Arg Leu Gly Ser Thr Val Phe Val Ala Asn Leu Asp Tyr Lys Val
                165                   170                   175
Gly
```

```
        <210> 203
        <211> 164
        <212> PRT
        <213> Homo sapien


        <400> 203
Met Arg Leu Ala Val Gly Ala Leu Leu Val Cys Ala Val Leu Gly Leu
1                   5                   10                   15
Cys Leu Ala Val Pro Asp Lys Thr Val Arg Trp Cys Ala Val Ser Glu
            20                   25                   30
His Glu Ala Thr Lys Cys Gln Ser Phe Arg Asp His Met Lys Ser Val
        35                   40                   45
Ile Pro Ser Asp Gly Pro Ser Val Ala Cys Val Lys Lys Ala Ser Tyr
    50                   55                   60
Leu Asp Cys Ile Arg Ala Ile Ala Ala Asn Glu Ala Asp Ala Val Thr
65                   70                   75                   80
Leu Asp Ala Gly Leu Val Tyr Asp Ala Tyr Leu Ala Pro Asn Asn Leu
                85                   90                   95
Lys Pro Val Val Ala Glu Phe Tyr Gly Ser Lys Glu Asp Pro Gln Thr
                100                  105                  110
Phe Tyr Tyr Ala Val Ala Val Val Lys Lys Asp Ser Gly Phe Gln Met
        115                  120                  125
Asn Gln Leu Arg Gly Lys Lys Ser Cys His Thr Gly Leu Gly Arg Ser
        130                  135                  140
Ala Gly Trp Asn Ile Pro Ile Gly Leu Leu Tyr Cys Asp Leu Pro Glu
145                  150                  155                  160
Pro Arg Lys Pro
```

```
        <210> 204
        <211> 241
        <212> PRT
        <213> Homo sapien


        <400> 204
Met Ser Gly Glu Ser Ala Arg Ser Leu Gly Lys Gly Ser Ala Pro Pro
1                   5                   10                   15
Gly Pro Val Pro Glu Gly Ser Ile Arg Ile Tyr Ser Met Arg Phe Cys
            20                   25                   30
Pro Phe Ala Glu Arg Thr Arg Leu Val Leu Lys Ala Lys Gly Ile Arg
        35                   40                   45
His Glu Val Ile Asn Ile Asn Leu Lys Asn Lys Pro Glu Trp Phe Phe
    50                   55                   60
Lys Lys Asn Pro Phe Gly Leu Val Pro Val Leu Glu Asn Ser Gln Gly
65                   70                   75                   80
```

```
Gln Leu Ile Tyr Glu Ser Ala Ile Thr Cys Glu Tyr Leu Asp Glu Ala
              85                    90                    95
Tyr Pro Gly Lys Lys Leu Leu Pro Asp Asp Pro Tyr Glu Lys Ala Cys
            100                 105                 110
Gln Lys Met Ile Leu Glu Leu Phe Ser Lys Val Pro Ser Leu Val Gly
        115                 120                 125
Ser Phe Ile Arg Ser Gln Asn Lys Glu Asp Tyr Asp Gly Leu Lys Glu
    130                     135                 140
Glu Phe Arg Lys Glu Phe Thr Lys Leu Glu Glu Val Leu Thr Asn Lys
145                 150                 155                 160
Lys Thr Thr Phe Phe Gly Gly Asn Ser Ile Ser Met Ile Asp Tyr Leu
                165                 170                 175
Ile Trp Pro Trp Phe Glu Arg Leu Glu Ala Met Lys Leu Asn Glu Cys
            180                 185                 190
Val Asp His Thr Pro Lys Leu Lys Leu Trp Met Ala Ala Met Lys Glu
        195                 200                 205
Asp Pro Thr Val Ser Ala Leu Leu Thr Ser Glu Lys Asp Trp Gln Gly
        210                 215                 220
Phe Leu Glu Leu Tyr Leu Gln Asn Ser Pro Glu Ala Cys Asp Tyr Gly
225                 230                 235                 240
Leu
```

```
        <210> 205
        <211> 160
        <212> PRT
        <213> Homo sapien


        <400> 205
Met Gln Ile Phe Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu
1               5                   10                  15
Val Glu Pro Ser Asp Thr Ile Glu Asn Val Lys Ala Lys Ile Gln Asp
            20                  25                  30
Lys Glu Gly Ile Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys
        35                  40                  45
Gln Leu Glu Asp Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu
    50                  55                  60
Ser Thr Leu His Leu Val Leu Arg Leu Arg Gly Gly Met Gln Ile Phe
65                  70                  75                  80
Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu Val Glu Pro Ser
                85                  90                  95
Asp Thr Ile Glu Asn Val Lys Ala Lys Ile Gln Asp Lys Glu Gly Ile
            100                 105                 110
Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys Gln Leu Glu Asp
        115                 120                 125
Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu Ser Thr Leu His
    130                 135                 140
Leu Val Leu Arg Leu Arg Gly Gly Met Gln Ile Phe Val Lys Thr Leu
145                 150                 155                 160
```

```
        <210> 206
        <211> 197
        <212> PRT
        <213> Homo sapien
```

```
      <400> 206
Thr Ser Pro Ser Glu Ala Cys Ala Pro Leu Leu Ile Ser Leu Ser Thr
1               5                   10              15
Leu Ile Tyr Asn Gly Ala Leu Pro Cys Gln Cys Asn Pro Gln Gly Ser
            20              25              30
Leu Ser Ser Glu Cys Asn Pro His Gly Gly Gln Cys Leu Cys Lys Pro
            35              40              45
Gly Val Val Gly Arg Arg Cys Asp Leu Cys Ala Pro Gly Tyr Tyr Gly
    50              55              60
Phe Gly Pro Thr Gly Cys Gln Gly Ala Cys Leu Gly Cys Arg Asp His
65              70              75              80
Thr Gly Gly Glu His Cys Glu Arg Cys Ile Ala Gly Phe His Gly Asp
            85              90              95
Pro Arg Leu Pro Tyr Gly Gly Gln Cys Arg Pro Cys Pro Cys Pro Glu
            100             105             110
Gly Pro Gly Ser Gln Arg His Phe Ala Thr Ser Cys His Gln Asp Glu
            115             120             125
Tyr Ser Gln Gln Ile Val Cys His Cys Arg Ala Gly Tyr Thr Gly Leu
    130             135             140
Arg Cys Glu Ala Cys Ala Pro Gly His Phe Gly Asp Pro Ser Arg Pro
145             150             155             160
Gly Gly Arg Cys Gln Leu Cys Glu Cys Ser Gly Asn Ile Asp Pro Met
            165             170             175
Asp Pro Asp Ala Cys Asp Pro His Thr Gly Gln Cys Leu Arg Cys Leu
            180             185             190
His His Thr Glu Gly
            195

      <210> 207
      <211> 175
      <212> PRT
      <213> Homo sapien

      <400> 207
Ile Ile Arg Gln Gln Gly Leu Ala Ser Tyr Asp Tyr Val Arg Arg Arg
1               5                   10              15
Leu Thr Ala Glu Asp Leu Phe Glu Ala Arg Ile Ile Ser Leu Glu Thr
            20              25              30
Tyr Asn Leu Leu Arg Glu Gly Thr Arg Ser Leu Arg Glu Ala Leu Glu
            35              40              45
Ala Glu Ser Ala Trp Cys Tyr Leu Tyr Gly Thr Gly Ser Val Ala Gly
    50              55              60
Val Tyr Leu Pro Gly Ser Arg Gln Thr Leu Ser Ile Tyr Gln Ala Leu
.65             70              75              80
Lys Lys Gly Leu Leu Ser Ala Glu Val Ala Arg Leu Leu Leu Glu Ala
            85              90              95
Gln Ala Ala Thr Gly Phe Leu Leu Asp Pro Val Lys Gly Glu Arg Leu
            100             105             110
Thr Val Asp Glu Ala Val Arg Lys Gly Leu Val Gly Pro Glu Leu His
            115             120             125
Asp Arg Leu Leu Ser Ala Glu Arg Ala Val Thr Gly Tyr Arg Asp Pro
    130             135             140
Tyr Thr Glu Gln Thr Ile Ser Leu Phe Gln Ala Met Lys Lys Glu Leu
145             150             155             160
Ile Pro Thr Glu Glu Ala Leu Arg Leu Trp Met Pro Ser Trp Pro
```

165   170   175

<210> 208
<211> 177
<212> PRT
<213> Homo sapien

<400> 208
Met Ala Ala Gly Val Glu Ala Ala Ala Glu Val Ala Ala Thr Glu Ile
1   5   10   15
Lys Met Glu Glu Glu Ser Gly Ala Pro Gly Val Pro Ser Gly Asn Gly
        20   25   30
Ala Pro Gly Pro Lys Gly Glu Gly Glu Arg Pro Ala Gln Asn Glu Lys
    35   40   45
Arg Lys Glu Lys Asn Ile Lys Arg Gly Gly Asn Arg Phe Glu Pro Tyr
    50   55   60
Ala Asn Pro Thr Lys Arg Tyr Arg Ala Phe Ile Thr Asn Ile Pro Phe
65   70   75   80
Asp Val Lys Trp Gln Ser Leu Lys Asp Leu Val Lys Glu Lys Val Gly
                85   90   95
Glu Val Thr Tyr Val Glu Leu Leu Met Asp Ala Glu Gly Lys Ser Arg
            100   105   110
Gly Cys Ala Val Val Glu Phe Lys Met Glu Glu Ser Met Lys Lys Ala
        115   120   125
Ala Glu Val Leu Asn Lys His Ser Leu Ser Gly Arg Pro Leu Lys Val
    130   135   140
Lys Glu Asp Pro Asp Gly Glu His Ala Arg Arg Ala Met Gln Lys Val
145   150   155   160
Met Ala Thr Thr Gly Gly Met Gly Met Gly Pro Gly Gly Pro Gly Met
                165   170   175
Ile

<210> 209
<211> 196
<212> PRT
<213> Homo sapien

<400> 209
Asp Leu Gln Asp Met Phe Ile Val His Thr Ile Glu Glu Ile Glu Gly
1   5   10   15
Leu Ile Ser Ala His Asp Gln Phe Lys Ser Thr Leu Pro Asp Ala Asp
            20   25   30
Arg Glu Arg Glu Ala Ile Leu Ala Ile His Lys Glu Ala Gln Arg Ile
        35   40   45
Ala Glu Ser Asn His Ile Lys Leu Ser Gly Ser Asn Pro Tyr Thr Thr
    50   55   60
Val Thr Pro Gln Ile Ile Asn Ser Lys Trp Glu Lys Val Gln Gln Leu
65   70   75   80
Val Pro Lys Arg Asp His Ala Leu Leu Glu Glu Gln Ser Lys Gln Gln
                85   90   95
Ser Asn Glu His Leu Arg Arg Gln Phe Ala Ser Gln Ala Asn Val Val
            100   105   110
Gly Pro Trp Ile Gln Thr Lys Met Glu Glu Ile Gly Arg Ile Ser Ile
        115   120   125

```
Glu Met Asn Gly Thr Leu Glu Asp Gln Leu Ser His Leu Lys Gln Tyr
    130                 135                 140
Glu Arg Ser Ile Val Asp Tyr Lys Pro Asn Leu Asp Leu Leu Glu Gln
145                 150                 155                 160
Gln His Gln Leu Ile Gln Glu Ala Leu Ile Phe Asp Asn Lys His Thr
                165                 170                 175
Asn Tyr Thr Met Glu His Ile Arg Val Gly Trp Glu Gln Leu Leu Thr
                180                 185                 190
Thr Ile Ala Arg
        195
```

```
        <210> 210
        <211> 156
        <212> PRT
        <213> Homo sapien

        <400> 210
Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val Ala Glu Gly Lys Glu
  1                 5                 10                  15
Val Leu Leu Leu Ala His Asn Leu Pro Gln Asn Arg Ile Gly Tyr Ser
                20                  25                  30
Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Ser Leu Ile Val Gly Tyr
        35                  40                  45
Val Ile Gly Thr Gln Gln Ala Thr Pro Gly Pro Ala Tyr Ser Gly Arg
    50                  55                  60
Glu Thr Ile Tyr Pro Asn Ala Ser Leu Leu Ile Gln Asn Val Thr Gln
65                  70                  75                  80
Asn Asp Thr Gly Phe Tyr Thr Leu Gln Val Ile Lys Ser Asp Leu Val
                85                  90                  95
Asn Glu Glu Ala Thr Gly Gln Phe His Val Tyr Pro Glu Leu Pro Lys
                100                 105                 110
Pro Ser Ile Ser Ser Asn Asn Ser Asn Pro Val Glu Asp Lys Asp Ala
            115                 120                 125
Val Ala Phe Thr Cys Glu Pro Glu Val Gln Asn Thr Thr Tyr Leu Trp
    130                 135                 140
Trp Val Asn Gly Gln Ser Leu Pro Val Ser Pro Lys
145                 150                 155
```

```
        <210> 211
        <211> 92
        <212> PRT
        <213> Homo sapien

        <400> 211
Met Glu Ser Pro Ser Ala Pro Pro His Arg Trp Cys Ile Pro Trp Gln
  1                 5                 10                  15
Arg Leu Leu Leu Thr Ala Ser Leu Leu Thr Phe Trp Asn Pro Pro Thr
                20                  25                  30
Thr Ala Lys Leu Thr Ile Glu Ser Thr Pro Phe Asn Val Ala Glu Gly
        35                  40                  45
Lys Glu Val Leu Leu Leu Val His Asn Leu Pro Gln His Leu Phe Gly
    50                  55                  60
Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Arg Gln Ile Ile
65                  70                  75                  80
Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr Pro Gly
```

```
<210> 212
<211> 142
<212> PRT
<213> Homo sapien

<400> 212
Glu Lys Gln Lys Asn Lys Glu Phe Ser Gln Thr Leu Glu Asn Glu Lys
1               5                   10                  15
Asn Thr Leu Leu Ser Gln Ile Ser Thr Lys Asp Gly Glu Leu Lys Met
            20                  25                  30
Leu Gln Glu Glu Val Thr Lys Met Asn Leu Leu Asn Gln Gln Ile Gln
        35                  40                  45
Glu Glu Leu Ser Arg Val Thr Lys Leu Lys Glu Thr Ala Glu Glu Glu
    50                  55                  60
Lys Asp Asp Leu Glu Glu Arg Leu Met Asn Gln Leu Ala Glu Leu Asn
65              70                  75                  80
Gly Ser Ile Gly Asn Tyr Cys Gln Asp Val Thr Asp Ala Gln Ile Lys
                85                  90                  95
Asn Glu Leu Leu Glu Ser Glu Met Lys Asn Leu Lys Lys Cys Val Ser
            100                 105                 110
Glu Leu Glu Glu Glu Lys Gln Gln Leu Val Lys Glu Lys Thr Lys Val
        115                 120                 125
Glu Ser Glu Ile Arg Lys Glu Tyr Leu Glu Lys Ile Gln Gly
    130                 135                 140

<210> 213
<211> 142
<212> PRT
<213> Homo sapien

<400> 213
Gly Gly Tyr Gly Gly Gly Tyr Gly Gly Val Leu Thr Ala Ser Asp Gly
1               5                   10                  15
Leu Leu Ala Gly Asn Glu Lys Leu Thr Met Gln Asn Leu Asn Asp Arg
            20                  25                  30
Leu Ala Ser Tyr Leu Asp Lys Val Arg Ala Leu Glu Ala Ala Asn Gly
        35                  40                  45
Glu Leu Glu Val Lys Ile Arg Asp Trp Tyr Gln Lys Gln Gly Pro Gly
    50                  55                  60
Pro Ser Arg Asp Tyr Ser His Tyr Tyr Thr Thr Ile Gln Asp Leu Arg
65              70                  75                  80
Asp Lys Ile Leu Gly Ala Thr Ile Glu Asn Ser Arg Ile Val Leu Gln
                85                  90                  95
Ile Asp Asn Ala Arg Leu Ala Ala Asp Asp Phe Arg Thr Lys Phe Glu
            100                 105                 110
Thr Glu Gln Ala Leu Arg Met Ser Val Glu Ala Asp Ile Asn Gly Leu
        115                 120                 125
Arg Arg Val Leu Asp Glu Leu Thr Leu Ala Arg Thr Asp Leu
    130                 135                 140

<210> 214
<211> 129
<212> PRT
```

<213> Homo sapien

<400> 214

```
Val Met Arg Val Asp Phe Asn Val Pro Met Lys Asn Asn Gln Ile Thr
1               5                   10                  15
Asn Asn Gln Arg Ile Lys Ala Ala Val Pro Ser Ile Lys Phe Cys Leu
            20                  25                  30
Asp Asn Gly Ala Lys Ser Val Val Leu Met Ser His Leu Gly Arg Pro
            35                  40                  45
Asp Gly Val Pro Met Pro Asp Lys Tyr Ser Leu Glu Pro Val Ala Val
        50                  55                  60
Glu Leu Arg Ser Leu Leu Gly Lys Asp Val Leu Phe Leu Lys Asp Cys
65                  70                  75                  80
Val Gly Pro Glu Val Glu Lys Ala Cys Ala Asn Pro Ala Ala Gly Ser
                85                  90                  95
Val Ile Leu Leu Glu Asn Leu Arg Phe His Val Glu Glu Glu Gly Lys
            100                 105                 110
Gly Lys Asp Ala Ser Gly Asn Lys Val Lys Ala Glu Pro Ala Lys Ile
            115                 120                 125
Glu
```

<210> 215
<211> 148
<212> PRT
<213> Homo sapien

<400> 215

```
Met Ala Thr Leu Lys Glu Lys Leu Ile Ala Pro Val Ala Glu Glu Glu
1               5                   10                  15
Ala Thr Val Pro Asn Asn Lys Ile Thr Val Val Gly Val Gly Gln Val
            20                  25                  30
Gly Met Ala Cys Ala Ile Ser Ile Leu Gly Lys Ser Leu Ala Asp Glu
            35                  40                  45
Leu Ala Leu Val Asp Val Leu Glu Asp Lys Leu Lys Gly Glu Met Met
        50                  55                  60
Asp Leu Gln His Gly Ser Leu Phe Leu Gln Thr Pro Lys Ile Val Ala
65                  70                  75                  80
Asp Lys Asp Tyr Ser Val Thr Ala Asn Ser Lys Ile Val Val Val Thr
                85                  90                  95
Ala Gly Val Arg Gln Gln Glu Gly Glu Ser Arg Leu Asn Leu Val Gln
            100                 105                 110
Arg Asn Val Asn Val Phe Lys Phe Ile Ile Pro Gln Ile Val Lys Tyr
            115                 120                 125
Ser Pro Asp Cys Ile Ile Ile Val Val Ser Asn Pro Val Asp Ile Leu
        130                 135                 140
Thr Tyr Val Thr
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<213> Homo sapien

<400> 216

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Gly Ser Pro Glu Arg Glu Glu Lys Val Leu Glu Asn Gly Glu Leu Thr
            20              25              30
Pro Pro Arg Arg Glu Glu Lys Ala Leu Glu Asn Gly Glu Leu Arg Ser
        35              40              45
Pro Glu Ala Gly Glu Lys Val Leu Val Asn Gly Gly Leu Thr Pro Pro
    50              55              60
Lys Ser Glu Asp Lys Val Ser Glu Asn Gly Gly Leu Arg Phe Pro Arg
65              70              75              80
Asn Thr Glu Arg Pro Pro Glu Thr Gly Pro Trp Arg Ala Pro Gly Pro
            85              90              95
Trp Glu Lys Thr Pro Glu Ser Trp Gly Pro Ala Pro Thr Ile Gly Glu
            100             105             110
Pro Ala Pro Glu Thr Ser Leu Glu Arg Ala Pro Ala Pro Ser Ala Val
        115             120             125
Val Ser Ser Arg Asn Gly Gly Glu Thr Ala Pro Gly Pro Leu Gly Pro
    130             135             140
Ala Pro Lys Asn Gly Thr Leu Glu Pro Gly Thr Glu Arg Arg Ala Pro
145             150             155             160
Glu Thr Gly Gly Ala Pro Arg Ala Pro Gly Ala Gly Arg Leu Asp Leu
            165             170             175
Gly Ser Gly Gly Arg Ala Pro Val Gly Thr Gly Thr Ala Pro Gly Gly
            180             185             190
Gly Pro Gly Ser Gly Val Asp Ala Lys Ala Gly Trp Val Asp Asn Thr
            195             200             205
Arg Pro Gln Pro Pro Pro Pro Leu Pro Pro Pro Pro Glu Ala Gln
    210             215             220
Pro Arg Arg Leu Glu Pro Ala Pro Pro Arg Ala Arg Pro Glu Val Ala
225             230             235             240
Pro Glu Gly Glu Pro Gly Ala Pro Asp Ser Arg Ala Gly Gly Asp Thr
            245             250             255
Ala Leu Ser Gly Asp Gly Asp Pro Pro Lys Pro Glu Arg Lys Gly Pro
            260             265             270
Glu Met Pro Arg Leu Phe Leu Asp Leu Gly Pro Pro Gln Gly Asn Ser
            275             280             285
Glu Gln Ile Lys Ala Arg Leu Ser Arg Leu Ser Leu Ala Leu Pro Pro
    290             295             300
Leu Thr Leu Thr Pro Phe Pro Gly Pro Gly Pro Arg Arg Pro Pro Trp
305             310             315             320
Glu Gly Ala Asp Ala Gly Ala Ala Gly Gly Glu Ala Gly Gly Ala Gly
            325             330             335
Ala Pro Gly Pro Ala Glu Glu Asp Gly Glu Asp Glu Asp Glu Asp Glu
            340             345             350
Glu Glu Asp Glu Glu Ala Ala Ala Pro Gly Ala Ala Ala Gly Pro Arg
            355             360             365
Gly Pro Gly Arg Ala Arg Ala Ala Pro Val Pro Val Val Val Ser Ser
    370             375             380
Ala Asp Ala Asp Ala Ala Arg Pro Leu Arg Gly Leu Leu Lys Ser Pro
385             390             395             400
Arg Gly Ala Asp Glu Pro Glu Asp Ser Glu Leu Glu Arg Lys Arg Lys
            405             410             415
Met Val Ser Phe His Gly Asp Val Thr Val Tyr Leu Phe Asp Gln Glu
            420             425             430
Thr Pro Thr Asn Glu Leu Ser Val Gln Ala Pro Pro Glu Gly Asp Thr
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                   435                      440                      445
        Asp Pro Ser Thr Pro Pro Ala Pro Pro Thr Pro Pro His Pro Ala Thr
            450                      455                      460
        Pro Gly Asp Gly Phe Pro Ser Asn Asp Ser Gly Phe Gly Gly Ser Phe
        465                      470                      475                      480
        Glu Trp Ala Glu Asp Phe Pro Leu Leu Pro Pro Pro Gly Pro Pro Leu
                             485                      490                      495
        Cys Phe Ser Arg Phe Ser Val Ser Pro Ala Leu Glu Thr Pro Gly Pro
                   500                      505                      510
        Pro Ala Arg Ala Pro Asp Ala Arg Pro Ala Gly Pro Val Glu Asn
                   515                      520                      525
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(51) International Patent Classification 6 :	A3	(11) International Publication Number:	WO 99/38973
C12N 15/12, A61K 38/17, C07K 14/47, 16/18, A61K 35/14		(43) International Publication Date:	5 August 1999 (05.08.99)

(54) Title: COMPOUNDS FOR THERAPY AND DIAGNOSIS OF LUNG CANCER AND METHODS FOR THEIR USE

(57) Abstract

Compounds and methods for treating lung cancer are provided. The inventive compounds include polypeptides containing at least a portion of a lung tumor protein. Vaccines and pharmaceutical compositions for immunotherapy of lung cancer comprising such polypeptides, or polynucleotides encoding such polypeptides, are also provided, together with polynucleotides for preparing the inventive polypeptides.

INTERNATIONAL SEARCH REPORT

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 C12N15/12   A61K38/17   C07K14/47   C07K16/18   A61K35/14

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
IPC 6 C12N C12Q A61K C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	WO 96 30389 A (MILLENIUM PHARMACEUTICALS, INC.; SHYJAN A.) 3 October 1996 see page 112 - page 127 ---	1-60
A	WO 96 02552 A (CYTOCLONYL PHARMACEUTICS, INC.; TORCZYNSKI R. ET AL.) 1 February 1996 see the whole document ---	1-60
A	YOU L ET AL.: "Identification of early growth response gene-1 (Egr-1) as a phorbol myristate-induced gene in lung cancer cells by differential mRNA display" AM. J. RESPIR. CELL MOL. BIOL., vol. 17, no. 5, November 1997, pages 617-624, XP002106654 see page 618, left-hand column, paragraph 3 ---	1,2,4-7

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[X] Further documents are listed in the continuation of box C.   [X] Patent family members are listed in annex.

* Special categories of cited documents :

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier document but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.

"&" document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
21 June 1999	22 10 1999

Name and mailing address of the ISA	Authorized officer
European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax (+31-70) 340-3016	CUPIDO, M

Form PCT/ISA/210 (second sheet) (July 1992)

INTERNATIONAL SEARCH REPORT

Box I   Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. [X] Claims Nos.:
   because they relate to subject matter not required to be searched by this Authority, namely:

   Remark: Although claims 16, 17, 24-26, 32, 33, 48-53 and 56-58 are
   directed to a method of treatment of the human/animal body
   the search has been carried out and based on the alleged
   effects of the composition.

2. [ ] Claims Nos.:
   because they relate to parts of the International Application that do not comply with the prescribed requirements to such
   an extent that no meaningful International Search can be carried out, specifically:

3. [ ] Claims Nos.:
   because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II   Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

   see FURTHER INFORMATION sheet

1. [ ] As all required additional search fees were timely paid by the applicant, this International Search Report covers all
   searchable claims.

2. [ ] As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment
   of any additional fee.

3. [ ] As only some of the required additional search fees were timely paid by the applicant, this International Search Report
   covers only those claims for which fees were paid, specifically claims Nos.:

4. [X] No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is
   restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

   see FURTHER INFORMATION sheet, subject 1.

Remark on Protest   [ ] The additional search fees were accompanied by the applicant's protest.

                           [ ] No protest accompanied the payment of additional search fees.

Form PCT/ISA/210 (continuation of first sheet (1)) (July 1998)

INTERNATIONAL SEARCH REPORT

Information on patent family members

Patent document cited in search report		Publication date	Patent family member(s)		Publication date
WO 9630389	A	03-10-1996	US	5633161 A	27-05-1997
			AU	708746 B	12-08-1999
			AU	5437896 A	16-10-1996
			CA	2216717 A	03-10-1996
			EP	0817792 A	14-01-1998
			US	5674739 A	07-10-1997
WO 9602552	A	01-02-1996	US	5589579 A	31-12-1996
			AU	700915 B	14-01-1999
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			EP	0804451 A	05-11-1997
			JP	10503087 T	24-03-1998
			US	5773579 A	30-06-1998